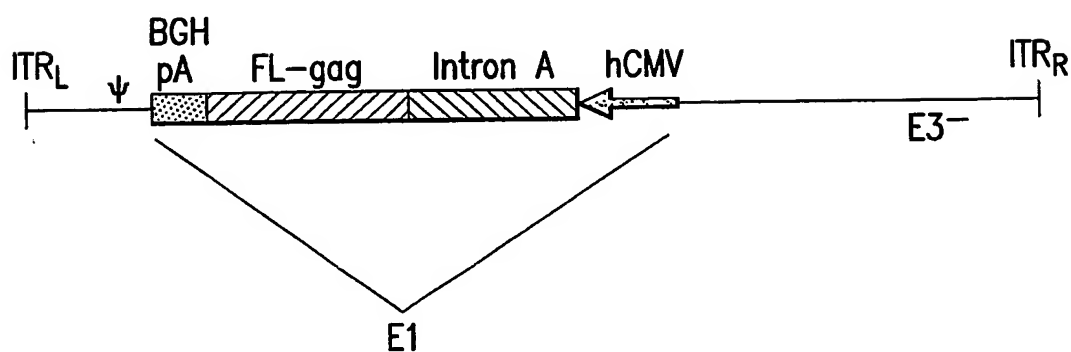


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ORIGINAL ADENOVECTOR CONSTRUCT:



ORIGINAL HIV-1 gag ADENOVECTOR.

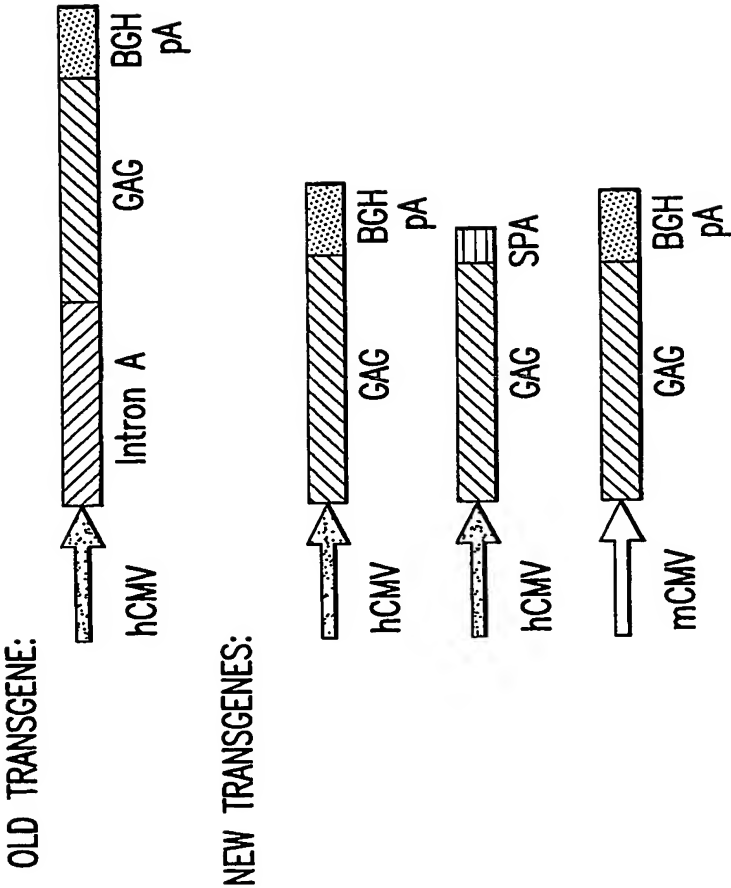
FIG.1

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Sequence of the open reading frame for FL-gag (human codon optimized)

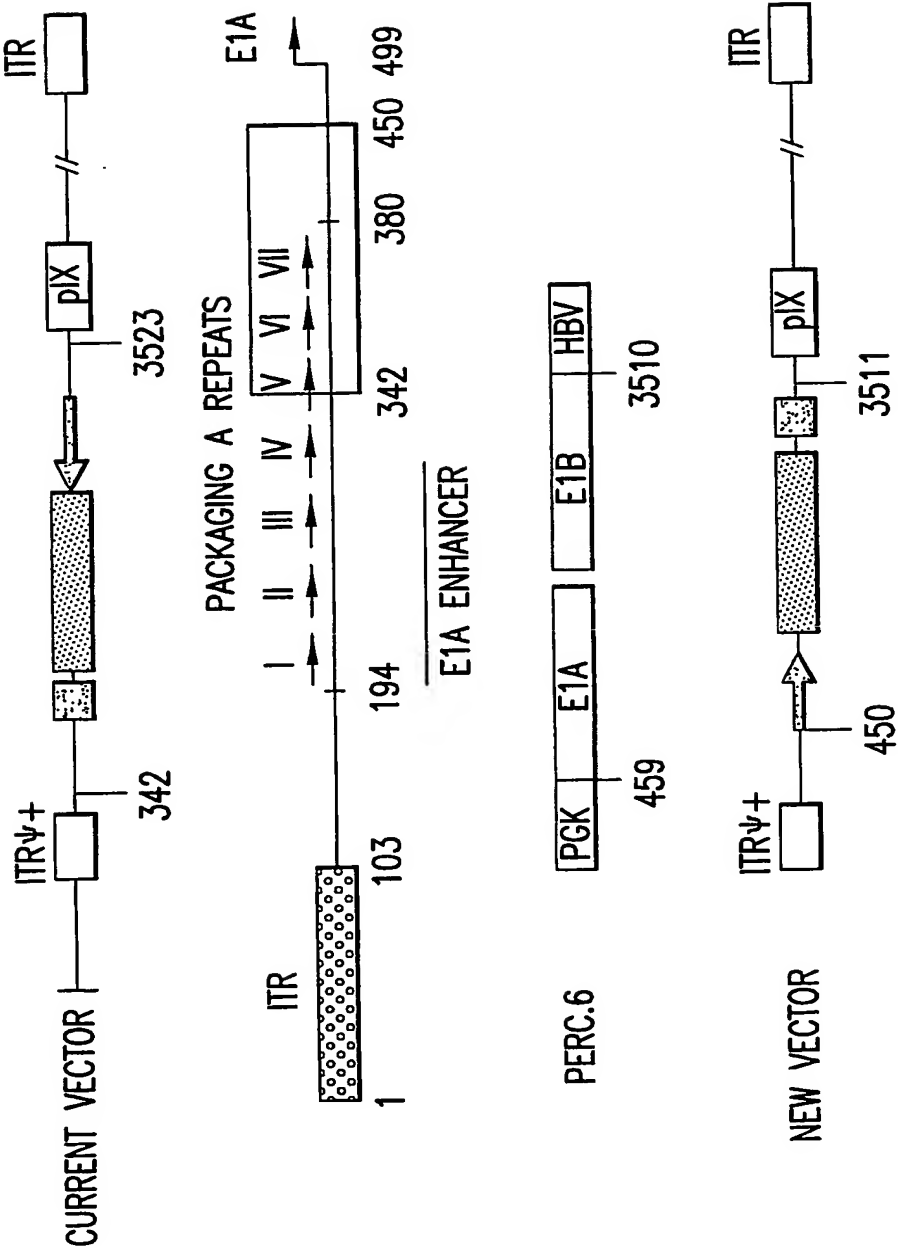
atgggtgctagggcttctgtgctgtctggtggtgagctggacaagtgggagaagatcaggctgaggcctggtg
caagaagaagtacaagctaaagcacattgtgtggcctccaggagctggagaggtttgctgtgaaccctggc
ctgctggagacctctgaggggtgcaggcagatcctgggccagctccagccctccctgcaaacaggctctgagg
agctgaggtccctgtacaacacagtggctaccctgtactgtgtgcaccagaagattgatgtgaaggacaccaag
gaggccctggagaagattgaggaggagcagaacaagtccaagaagaaggcccagcaggctgctgctggc
acaggcaactccagccagggtgtcccagaactacccattgtgcagaacctccaggggccagatggtgcaccag
gccatctccccccggaccctgaatgcctgggtgaaggtggtggaggagaaggccttctcccctgaggtgatccc
catgttctctgcccctgtctgaggggtgccacccccaggacctgaacaccatgctgaacacagtggggggccatc
aggctgccatgcagatgctgaaggagacctcaatgaggaggctgctgagtgggacaggctgcatcctgtgc
acgctggccccattgcccccgccagatgaggggagcccaggggctctgacattgctggcaccacctccacct
ccaggagcagattggctggatgaccaacaaccccccatccctgtgggggaaatctacaagaggtggatcat
cctgggcctgaacaagattgtgaggatgtactccccacctccatcctggacatcaggcaggggcccaaggag
cccttcagggactatgtggacaggttctacaagacctgagggtgagcaggcctccaggagggtgaagaact
ggatgacagagaccctgctggtgcagaatgccaacctgactgcaagaccatcctgaaggccctgggcccctg
ctgccacctggaggagatgatgacagcctgccaggggtggggggccctggtcacaaggccagggtgctg
gctgaggccatgtcccagggtgaccaactccgccaccatcatgatgcagaggggcaacttcaggaaccagag
gaagacagtgaagtgttcaactgtggcaaggtgggccacattgccaagaactgtagggccccccaggaaga
agggtgctggaagtgtggcaaggaggggccaccagatgaaggactgcaatgagaggcaggccaacttcctg
ggcaaaatctggccctcccacaagggcaggcctggcaacttcctccagtccaggcctgagcccacagcccct
cccaggagtccttcagggttggggaggagaagaccacccccagccagaagcaggagcccattgacaagg
agctgtacccccctggcctccctgaggtccctgtttggcaacgacctcctcccagtaaaataaagcccgggca
gat

FIG.2



DIAGRAMMATIC REPRESENTATION OF THE ORIGINAL HIV-1 GAG TRANSGENE AND THE SERIES OF NEW TRANSGENE CONSTRUCTIONS.

FIG.3



MODIFICATIONS MADE TO THE CURRENT ADENOVECTOR BACKBONE IN THE GENERATION OF THE NEW VECTOR.

FIG.4

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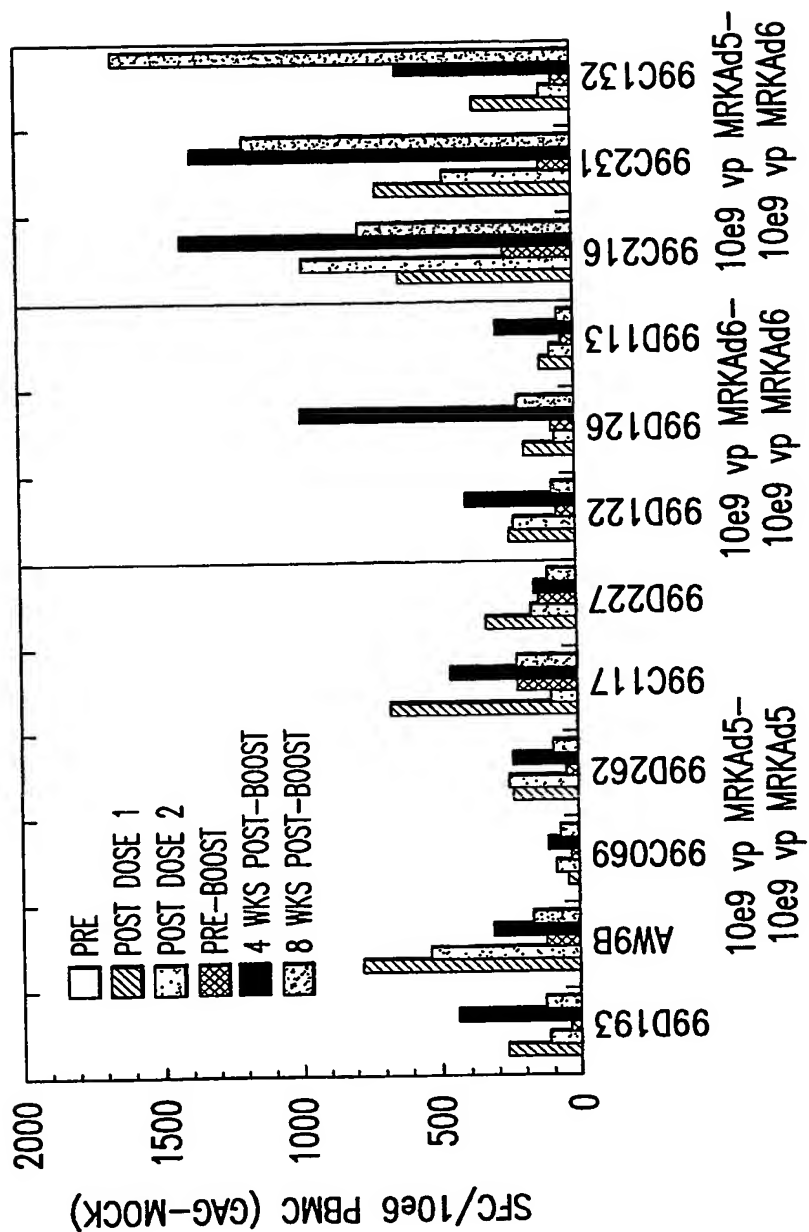


FIG.5

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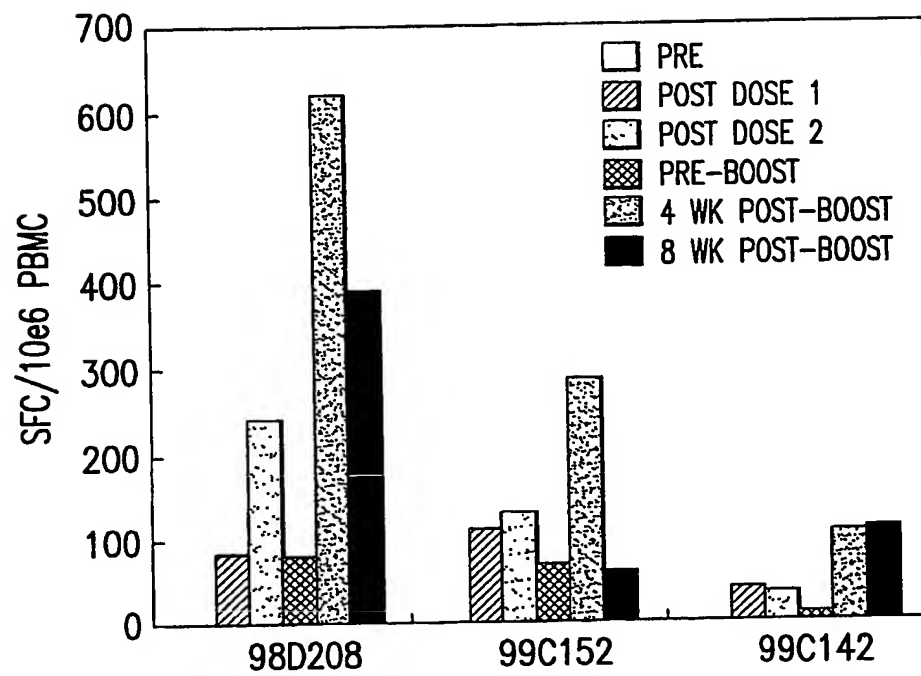


FIG.6

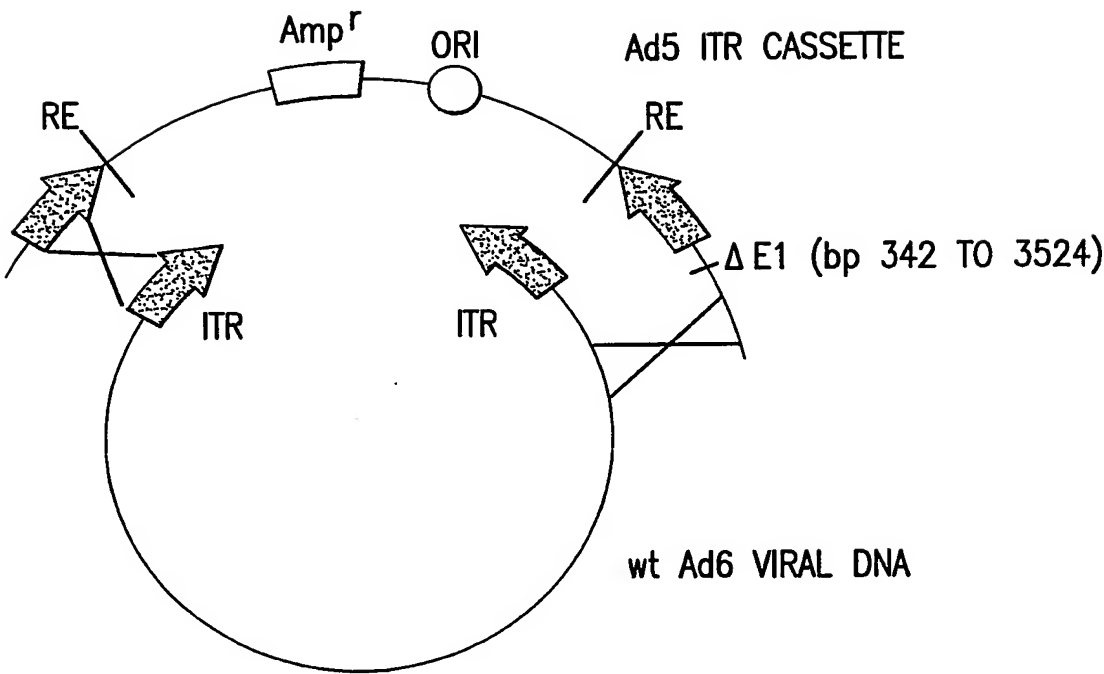


FIG.7

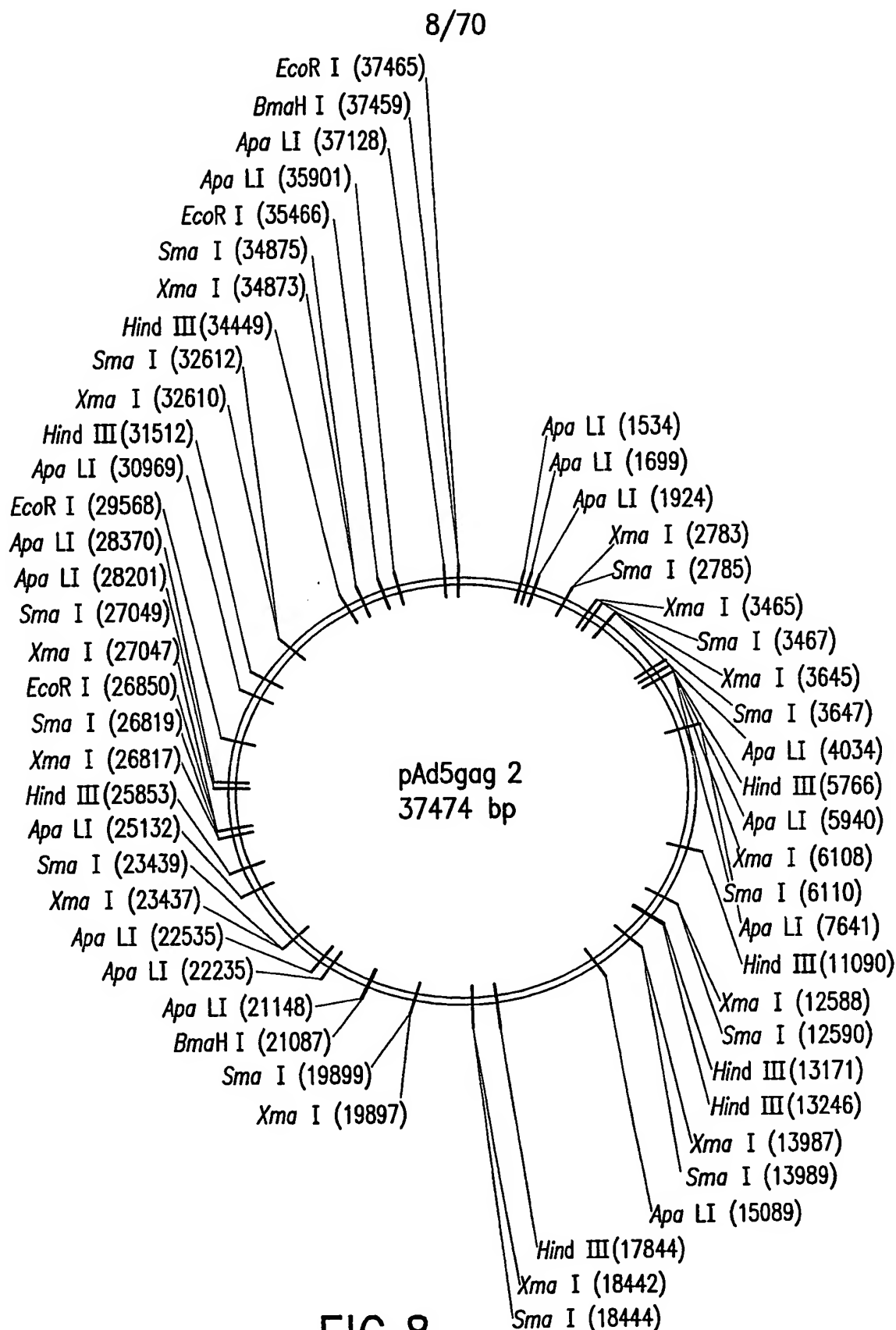


FIG.8

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1  TTCTTAATTA ACATCATCAA TAATATACCT TATTTTGGAT TGAAGCCAAT
   AAGAATTAAT TGTAGTAGTT ATTATATGGA ATAAACCTA ACTTCGGTTA

51  ATGATAATGA GGGGGTGGAG TTTGTGACGT GGCGCGGGGC GTGGGAACGG
   TACTATTACT CCCCACCTC AAACACTGCA CCGCGCCCCG CACCCTTGCC

101 GGCGGGTGAC GTAGTAGTGT GGCGGAAGTG TGATGTTGCA AGTGTGGCGG
   CCGCCCCTG CATCATCACA CCGCCTTCAC ACTACAACGT TCACACCGCC

151 AACACATGTA AGCGACGGAT GTGGCAAAAG TGACGTTTTT GGTGTGCGCC
   TTGTGTACAT TCGCTGCCTA CACCGTTTTT ACTGCAAAAA CCACACGCGG

201 GGTGTACACA GGAAGTGACA ATTTTCGCGC GGTTTTAGGC GGATGTTGTA
   CCACATGTGT CCTTCACTGT TAAAAGCGCG CCAAATCCG CCTACAACAT

251 GTAAATTTGG GCGTAACCGA GTAAGATTTG GCCATTTTCG CGGGAAACT
   CATTTAAACC CGCATTGGCT CATTCTAAAC CGGTAAAAGC GCCCTTTTGA

301 GAATAAGAGG AAGTGAAATC TGAATAATTT TGTGTTACTC ATAGCGCGTA
   CTTATTCTCC TTCCTTTAG ACTTATTAA ACACAATGAG TATCGCGCAT

351 ATATTTGTCT AGGGCCGCGG GGACTTTGAC CGTTTACGTG GAGACTCGCC
   TATAAACAGA TCCCGGCGCC CCTGAAACTG GCAAATGCAC CTCTGAGCGG

401 CAGGTGTTTT TCTCAGGTGT TTTCCGCGTT CCGGGTCAAA GTTGGCGTTT
   GTCCACAAAA AGAGTCCACA AAAGGCGCAA GGCCAGTTT CAACCGCAAA

451 TATTATTATA GCGGCGCGCG ATCCATTGCA TACGTTGTAT CCATATCATA
   ATAATAATAT CCGCCGGCGC TAGGTAACGT ATGCAACATA GGTATAGTAT

501 ATATGTACAT TTATATTGGC TCATGTCCAA CATTACCGCC ATGTTGACAT
   TATACATGTA AATATAACCG AGTACAGGTT GTAATGGCGG TACAACTGTA

551 TGATTATTGA CTAGTTATTA ATAGTAATCA ATTACGGGGT CATTAGTTCA
   ACTAATAACT GATCAATAAT TATCATTAGT TAATGCCCCA GTAATCAAGT

601 TAGCCCATAT ATGGAGTTCC GCGTTACATA ACTTACGGTA AATGGCCCGC
   ATCGGGTATA TACCTCAAGG CGCAATGTAT TGAATGCCAT TTACCGGGCG

651 CTGGCTGACC GCCCAACGAC CCCC GCCCAT TGACGTCAAT AATGACGTAT
   GACCGACTGG CGGGTTGCTG GGGGCGGGTA ACTGCAGTTA TTA CTGCATA

701 GTTCCCATAG TAACGCCAAT AGGGACTTTC CATTGACGTC AATGGGTGGA
   CAAGGGTATC ATTGCGGTTA TCCCTGAAAG GTA ACTGCAG TTA CCGACCT

751 GTATTTACGG TAAACTGCCC ACTTGGCAGT ACATCAAGTG TATCATATGC
   CATAAATGCC ATTTGACGGG TGAACCGTCA TG TAGTTCAC ATAGTATACG

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FIG.9A-1

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|      |             |             |              |             |            |
|------|-------------|-------------|--------------|-------------|------------|
| 801  | CAAGTACGCC  | CCCTATTGAC  | GTCAATGACG   | GTAAATGGCC  | CGCCTGGCAT |
|      | GTTCATGCGG  | GGGATAACTG  | CAGTTACTGC   | CATTTACCGG  | GCGGACCGTA |
| 851  | TATGCCCAGT  | ACATGACCTT  | ATGGGACTTT   | CCTACTTGGC  | AGTACATCTA |
|      | ATACGGGTCA  | TGTACTGGAA  | TACCCTGAAA   | GGATGAACCG  | TCATGTAGAT |
| 901  | CGTATTAGTC  | ATCGCTATTA  | CCATGGTGAT   | GCGGTTTTGG  | CAGTACATCA |
|      | GCATAATCAG  | TAGCGATAAT  | GGTACCACTA   | CGCCAAAACC  | GTCATGTAGT |
| 951  | ATGGGCGTGG  | ATAGCGGTTT  | GA CTCACGGG  | GATTTCCAAG  | TCTCCACCCC |
|      | TACCCGCACC  | TATCGCCAAA  | CTGAGTGCCC   | CTAAAGGTTT  | AGAGGTGGGG |
| 1001 | ATTGACGTCA  | ATGGGAGTTT  | GTTTTGGCAC   | CAAATCAAC   | GGGACTTTCC |
|      | TA ACTGCAGT | TACCCTCAAA  | CAA AACC GTG | GTTTTAGTTG  | CCCTGAAAGG |
| 1051 | AAAATGTCGT  | AACA ACTCCG | CCCCATTGAC   | GCAAATGGGC  | GGTAGGCGTG |
|      | TTTTACAGCA  | TTGTTGAGGC  | GGGGTAACTG   | CGTTTACCCG  | CCATCCGCAC |
| 1101 | TACGGTGGGA  | GGTCTATATA  | AGCAGAGCTC   | GTTTAGTGAA  | CCGTCAGATC |
|      | ATGCCACCCT  | CCAGATATAT  | TCGTCTCGAG   | CAAATCACTT  | GGCAGTCTAG |
| 1151 | GCCTGGAGAC  | GCCATCCACG  | CTGTTTTGAC   | CTCCATAGAA  | GACACCGGGA |
|      | CGGACCTCTG  | CGGTAGGTGC  | GACAAA ACTG  | GAGGTATCTT  | CTGTGGCCCT |
| 1201 | CCGATCCAGC  | CTCCGCGGCC  | GGGAACGGTG   | CATTGGAACG  | CGGATTCCCC |
|      | GGCTAGGTCTG | GAGGCGCCGG  | CCCTTGCCAC   | GTAACCTTGC  | GCCTAAGGGG |
| 1251 | GTGCCAAGAG  | TGAGATCTAC  | CATGGGTGCT   | AGGGCTTCTG  | TGCTGTCTGG |
|      | CACGGTTCTC  | ACTCTAGATG  | GTACCCACGA   | TCCCGAAGAC  | ACGACAGACC |
| 1301 | TGGTGAGCTG  | GACAAGTGGG  | AGAAGATCAG   | GCTGAGGCCT  | GGTGGAAGA  |
|      | ACCACTCGAC  | CTGTTCACCC  | TCTTCTAGTC   | CGACTCCGGA  | CCACCGTTCT |
| 1351 | AGAAGTACAA  | GCTAAAGCAC  | ATTGTGTGGG   | CCTCCAGGGA  | GCTGGAGAGG |
|      | TCTTCATGTT  | CGATTTCTGT  | TAACACACCC   | GGAGGTCCCT  | CGACCTCTCC |
| 1401 | TTTGCTGTGA  | ACCCTGGCCT  | GCTGGAGACC   | TCTGAGGGGT  | GCAGGCAGAT |
|      | AAACGACACT  | TGGGACCGGA  | CGACCTCTGG   | AGACTCCCCA  | CGTCCGTCTA |
| 1451 | CCTGGGCCAG  | CTCCAGCCCT  | CCCTGCAAAC   | AGGCTCTGAG  | GAGCTGAGGT |
|      | GGACCCGGTC  | GAGGTCGGGA  | GGGACGTTTG   | TCCGAGACTC  | CTCGACTCCA |
| 1501 | CCCTGTACAA  | CACAGTGGCT  | ACCCTGTACT   | GTGTGCACCA  | GAAGATTGAT |
|      | GGGACATGTT  | GTGTCACCGA  | TGGGACATGA   | CACACGTGGT  | CTTCTAACTA |
| 1551 | GTGAAGGACA  | CCAAGGAGGC  | CCTGGAGAAG   | ATTGAGGAGG  | AGCAGAACAA |
|      | CACTTCCTGT  | GGTTCCTCCG  | GGACCTCTTC   | TA ACTCCTCC | TCGTCTTGTT |
| 1601 | GTCCAAGAAG  | AAGGCCCAGC  | AGGCTGCTGC   | TGGCACAGGC  | AACTCCAGCC |
|      | CAGGTTCTTC  | TTCCGGGTCTG | TCCGACGACG   | ACCGTGTCCG  | TTGAGGTCGG |

FIG.9A-2

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1651 AGGTGTCCCA GAACTACCCC ATTGTGCAGA ACCTCCAGGG CCAGATGGTG  
 TCCACAGGGT CTTGATGGGG TAACACGTCT TGGAGGTCCC GGTCTACCAC  
 1701 CACCAGGCCA TCTCCCCCG GACCCTGAAT GCCTGGGTGA AGGTGGTGGA  
 GTGGTCCGGT AGAGGGGGGC CTGGGACTTA CGGACCCACT TCCACCACCT  
 1751 GGAGAAGGCC TTCTCCCCTG AGGTGATCCC CATGTTCTCT GCCCTGTCTG  
 CCTCTTCCGG AAGAGGGGAC TCCACTAGGG GTACAAGAGA CGGGACAGAC  
 1801 AGGGTGCCAC CCCCAGGAC CTGAACACCA TGCTGAACAC AGTGGGGGGC  
 TCCCACGGTG GGGGGTCCTG GACTTGTGGT ACGACTTGTG TCACCCCCCG  
 1851 CATCAGGCTG CCATGCAGAT GCTGAAGGAG ACCATCAATG AGGAGGCTGC  
 GTAGTCCGAC GGTACGTCTA CGACTTCCTC TGGTAGTTAC TCCTCCGACG  
 1901 TGAGTGGGAC AGGCTGCATC CTGTGCACGC TGGCCCCATT GCCCCCGGCC  
 ACTCACCCCTG TCCGACGTAG GACACGTGCG ACCGGGGTAA CGGGGGCCGG  
 1951 AGATGAGGGA GCCCAGGGGC TCTGACATTG CTGGCACCAC CTCCACCCTC  
 TCTACTCCCT CGGGTCCCCG AGACTGTAAC GACCGTGGTG GAGGTGGGAG  
 2001 CAGGAGCAGA TTGGCTGGAT GACCAACAAC CCCCCATCC CTGTGGGGGA  
 GTCCTCGTCT AACCGACCTA CTGGTTGTTG GGGGGGTAGG GACACCCCCT  
 2051 AATCTACAAG AGGTGGATCA TCCTGGGCCT GAACAAGATT GTGAGGATGT  
 TTAGATGTTT TCCACCTAGT AGGACCCGGA CTTGTTCTAA CACTCCTACA  
 2101 ACTCCCCCAC CTCCATCCTG GACATCAGGC AGGGCCCCAA GGAGCCCTTC  
 TGAGGGGGTG GAGGTAGGAC CTGTAGTCCG TCCCGGGGTT CCTCGGGAAG  
 2151 AGGGACTATG TGGACAGGTT CTACAAGACC CTGAGGGCTG AGCAGGCCTC  
 TCCCTGATAC ACCTGTCCAA GATGTTCTGG GACTCCCGAC TCGTCCGGAG  
 2201 CCAGGAGGTG AAGAACTGGA TGACAGAGAC CCTGCTGGTG CAGAATGCCA  
 GGTCTCTCAC TTCTTGACCT ACTGTCTCTG GGACGACCAC GTCTTACGGT  
 2251 ACCCTGACTG CAAGACCATC CTGAAGGCC TGGGCCCTGC TGCCACCCTG  
 TGGGACTGAC GTTCTGGTAG GACTTCCGGG ACCCGGGACG ACGGTGGGAC  
 2301 GAGGAGATGA TGACAGCCTG CCAGGGGGTG GGGGGCCCTG GTCACAAGGC  
 CTCCTCTACT ACTGTCGGAC GGTCCCCAC CCCCAGGAC CAGTGTTCCG  
 2351 CAGGGTGCTG GCTGAGGCCA TGTCCCAGGT GACCAACTCC GCCACCATCA  
 GTCCACGAC CGACTCCGGT ACAGGGTCCA CTGTTGAGG CGGTGGTAGT  
 2401 TGATGCAGAG GGGCAACTTC AGGAACCAGA GGAAGACAGT GAAGTGCTTC  
 ACTACGTCTC CCCGTTGAAG TCCTTGGTCT CTTCTGTCA CTTCACGAAG  
 2451 AACTGTGGCA AGGTGGGCCA CATTGCCAAG AACTGTAGGG CCCCAGGAA  
 TTGACACCGT TCCACCCGGT GTAACGGTTC TTGACATCCC GGGGGTCCTT

FIG.9A-3

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2501 GAAGGGCTGC TGGAA GTGTG GCAAGGAGGG CCACCAGATG AAGGACTGCA  
 CTTCCCGACG ACCTTCACAC CGTTCCTCCC GGTGGTCTAC TTCCTGACGT  
 2551 ATGAGAGGCA GGCCAACTTC CTGGGCAAAA TCTGGCCCTC CCACAAGGGC  
 TACTCTCCGT CCGGTTGAAG GACCCGTTTT AGACCGGGAG GGTGTTCCCG  
 2601 AGGCCTGGCA ACTTCCTCCA GTCCAGGCCT GAGCCCACAG CCCCTCCCGA  
 TCCGGACCGT TGAAGGAGGT CAGGTCCGGA CTCGGGTGTC GGGGAGGGCT  
 2651 GGAGTCCTTC AGGTTTGGGG AGGAGAAGAC CACCCCCAGC CAGAAGCAGG  
 CCTCAGGAAG TCCAAACCCC TCCTCTTCTG GTGGGGGTGTC GTCTTCGTCC  
 2701 AGCCCATTGA CAAGGAGCTG TACCCCTGG CCTCCCTGAG GTCCCTGTTT  
 TCGGGTAACT GTTCCTCGAC ATGGGGGACC GGAGGGACTC CAGGGACAAA  
 2751 GGCAACGACC CCTCCTCCCA GTAAAATAAA GCCCGGGCAG ATCTGCTGTG  
 CCGTTGCTGG GGAGGAGGGT CATTTTATTT CGGGCCCGTC TAGACGACAC  
 2801 CCTTCTAGTT GCCAGCCATC TGTTGTTTGC CCCTCCCCCG TGCTTTCCTT  
 GGAAGATCAA CGGTCGGTAG ACAACAAACG GGGAGGGGGC ACGGAAGGAA  
 2851 GACCCTGGAA GGTGCCACTC CCACTGTCCT TTCCTAATAA AATGAGGAAA  
 CTGGGACCTT CCACGGTGAG GGTGACAGGA AAGGATTATT TTA CTCTTT  
 2901 TTGCATCGCA TTGTCTGAGT AGGTGTCATT CTATTCTGGG GGGTGGGGTG  
 AACGTAGCGT AACAGACTCA TCCACAGTAA GATAAGACCC CCCACCCAC  
 2951 GGGCAGGACA GCAAGGGGGA GGATTGGGAA GACAATAGCA GGCATGCTGG  
 CCCGTCCTGT CGTTCCCCCT CTAACCCTT CTGTTATCGT CCGTACGACC  
 3001 GGATGCGGTG GGCTCTATGG CCGATCGGCG CGCCGTA CTG AAATGTGTGG  
 CCTACGCCAC CCGAGATACC GGCTAGCCGC GCGGCATGAC TTTACACACC  
 3051 GCGTGGCTTA AGGGTGGGAA AGAATATATA AGGTGGGGGT CTTATGTAGT  
 CGCACC GAAT TCCCACCCTT TCTTATATAT TCCACCCCA GAATACATCA  
 3101 TTTGTATCTG TTTTGCAGCA GCCGCCGCCG CCATGAGCAC CAACTCGTTT  
 AAACATAGAC AAAACGTCGT CGGCGGCGGC GGTACTCGTG GTTGAGCAAA  
 3151 GATGGAAGCA TTGTGAGCTC ATATTTGACA ACGCGCATGC CCCCATGGGC  
 CTACCTTCGT AACACTCGAG TATAAACTGT TGC GCGTACG GGGGTACCCG  
 3201 CGGGGTGCGT CAGAATGTGA TGGGCTCCAG CATTGATGGT CGCCCCGTCC  
 GCCCCACGCA GTCTTACACT ACCCGAGGTC GTA ACTACCA GCGGGGACAGG  
 3251 TGCCCGCAAA CTCTACTACC TTGACCTACG AGACCGTGTC TGG AACGCCG  
 ACGGGCGTTT GAGATGATGG AACTGGATGC TCTGGCACAG ACCTTGCGGC  
 3301 TTGGAGACTG CAGCCTCCGC CGCCGCTTCA GCCGCTGCAG CCACCGCCCCG  
 AACCTCTGAC GTCGGAGGCG GCGGCGAAGT CGGCGACGTC GGTGGCGGGC

FIG.9A-4

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3351 CGGGATTGTG ACTGACTTTG CTTTCCTGAG CCCGCTTGCA AACAGTGCAG  
 GCCCTAACAC TGA CTGAAAC GAAAGGACTC GGGCGAACGT TTGTCACGTC  
 3401 CTTCCCGTTC ATCCGCCCCG GATGACAAGT TGACGGCTCT TTTGGCACAA  
 GAAGGGCAAG TAGGCGGGCG C TACTGTTCA ACTGCCGAGA AAACCGTGTT  
 3451 TTGGATTCTT TGACCCGGGA ACTTAATGTC GTTTCTCAGC AGCTGTTGGA  
 AACCTAAGAA ACTGGGCCCT TGAATTACAG CAAAGAGTCG TCGACAACCT  
 3501 TCTGCGCCAG CAGGTTTCTG CCCTGAAGGC TTCCTCCCCT CCCAATGCGG  
 AGACGCGGTC GTCCAAAGAC GGGACTTCCG AAGGAGGGGA GGGTTACGCC  
 3551 TTTAAACAT AAATAAAAA CCAGACTCTG TTTGGATTG GATCAAGCAA  
 AAATTTTGTA TTTATTTTT GGTCTGAGAC AAACCTAAAC CTAGTTCGTT  
 3601 GTGTCTTGCT GTCTTTATTT AGGGGTTTTG CGCGCGCGGT AGGCCCGGGA  
 CACAGAACGA CAGAAATAAA TCCCCAAAAC GCGCGCGCCA TCCGGGCCCT  
 3651 CCAGCGGTCT CGGTCGTTGA GGGTCCTGTG TATTTTTTCC AGGACGTGGT  
 GGTGCGCCAGA GCCAGCAACT CCCAGGACAC ATAAAAAAGG TCCTGCACCA  
 3701 AAAGGTGACT CTGGATGTTT AGATACATGG GCATAAGCCC GTCTCTGGGG  
 TTTCCACTGA GACCTACAAG TCTATGTACC CGTATTCGGG CAGAGACCCC  
 3751 TGGAGGTAGC ACCACTGCAG AGCTTCATGC TCGGGGGTGG TGTGTAGAT  
 ACCTCCATCG TGGTGACGTC TCGAAGTACG ACGCCCCACC ACAACATCTA  
 3801 GATCCAGTCG TAGCAGGAGC GCTGGGCGTG GTGCCTAAAA ATGTCTTTCA  
 CTAGGTCAGC ATCGTCCTCG CGACCCGCAC CACGGATTTT TACAGAAAGT  
 3851 GTAGCAAGCT GATTGCCAGG GGCAGGCCCT TGGTGTAAGT GTTTACAAAG  
 CATCGTTTGA CTAACGGTCC CCGTCCGGGA ACCACATTCA CAAATGTTTC  
 3901 CGGTTAAGCT GGGATGGGTG CATACTGTTG GATATGAGAT GCATCTTGGA  
 GCCAATTCGA CCTACCCAC GTATGCACCC CTATACTCTA CGTAGAACCT  
 3951 CTGTATTTTT AGGTTGGCTA TGTTCCCAGC CATATCCCTC CGGGGATTCA  
 GACATAAAAA TCCAACCGAT ACAAGGGTCG GTATAGGGAG GCCCCTAAGT  
 4001 TGTGTGTCAG AACCACCAGC ACAGTGTATC CGGTGCACTT GGGAAATTTG  
 ACAACACGTC TTGGTGGTCG TGTCACATAG GCCACGTGAA CCCTTTAAAC  
 4051 TCATGTAGCT TAGAAGGAAA TCGTGGAAG AACTTGGAGA CGCCCTTGTTG  
 AGTACATCGA ATCTTCCTTT ACGCACCTTC TTGAACCTCT GCGGGAACAC  
 4101 ACCTCCAAGA TTTTCCATGC ATTCGTCCAT AATGATGGCA ATGGGCCCCAC  
 TGGAGGTTCT AAAAGGTACG TAAGCAGGTA T TACTACCGT TACCCGGGTG  
 4151 GGGCGGCGGC CTGGGCGAAG ATATTTCTGG GATCACTAAC GTCATAGTTG  
 CCCGCCGCCG GACCCGCTTC TATAAAGACC CTAGTGATTG CAGTATCAAC

FIG.9A-5

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4201 TGTTCCAGGA TGAGATCGTC ATAGGCCATT TTTACAAAGC GCGGGCGGAG  
 ACAAGGTCCT ACTCTAGCAG TATCCGGTAA AAATGTTTCG CGCCCGCCTC  
 4251 GGTGCCAGAC TGCGGTATAA TGGTTCCATC CGGCCCAGGG GCGTAGTTAC  
 CCACGGTCTG ACGCCATATT ACCAAGGTAG GCCGGGTCCC CGCATCAATG  
 4301 CCTCACAGAT TTGCATTTCC CACGCTTTGA GTTCAGATGG GGGGATCATG  
 GGAGTGTCTA AACGTAAAGG GTGCGAAACT CAAGTCTACC CCCCTAGTAC  
 4351 TCTACCTGCG GGGCGATGAA GAAAACGGTT TCCGGGGTAG GGGAGATCAG  
 AGATGGACGC CCCGCTACTT CTTTGGCAA AGGCCCATC CCCTCTAGTC  
 4401 CTGGGAAGAA AGCAGGTTCC TGAGCAGCTG CGACTTACCG CAGCCGGTGG  
 GACCTTCTT TCGTCCAAGG ACTCGTCGAC GCTGAATGGC GTCGGCCACC  
 4451 GCCCGTAAAT CACACCTATT ACCGGCTGCA ACTGGTAGTT AAGAGAGCTG  
 CGGGCATTTA GTGTGGATAA TGGCCGACGT TGACCATCAA TTCTCTCGAC  
 4501 CAGCTGCCGT CATCCCTGAG CAGGGGGGCC ACTTCGTTAA GCATGTCCCT  
 GTCGACGGCA GTAGGGACTC GTCCCCCGG TGAAGCAATT CGTACAGGGA  
 4551 GACTCGCATG TTTTCCCTGA CCAAATCCGC CAGAAGGCGC TCGCCGCCA  
 CTGAGCGTAC AAAAGGGACT GGTTTAGGCG GTCTTCCGCG AGCGGCGGGT  
 4601 GCGATAGCAG TTCTTGCAAG GAAGCAAAGT TTTTCAACGG TTTGAGACCG  
 CGCTATCGTC AAGAACGTTT CTTTCGTTTCA AAAAGTTGCC AAACCTCTGGC  
 4651 TCCGCCGTAG GCATGCTTTT GAGCGTTTGA CCAAGCAGTT CCAGGCGGTG  
 AGGCGGCATC CGTACGAAAA CTCGCAAACT GGTTCGTCAA GGTCCGCCAG  
 4701 CCACAGCTCG GTCACCTGCT CTACGGCATC TCGATCCAGC ATATCTCCTC  
 GGTGTGAGC CAGTGGACGA GATGCCGTAG AGCTAGGTCG TATAGAGGAG  
 4751 GTTTCGCGGG TTGGGGCGGC TTTTCGCTGTA CGGCAGTAGT CCGTGCTCGT  
 CAAAGCGCCC AACCCCGCCG AAAGCGACAT GCCGTCATCA GCCACGAGCA  
 4801 CCAGACGGGC CAGGGTCATG TCTTTCCACG GGCGCAGGGT CCTCGTCAGC  
 GGTCTGCCCG GTCCCAGTAC AGAAAGGTGC CCGCGTCCCA GGAGCAGTCG  
 4851 GTAGTCTGGG TCACGGTGAA GGGGTGCGCT CCGGGCTGCG CGCTGGCCAG  
 CATCAGACCC AGTGCCACTT CCCCACGCGA GGCCCGACGC GCGACCGGTC  
 4901 GGTGCGCTTG AGGCTGGTCC TGCTGGTGCT GAAGCGCTGC CCGTCTTCGC  
 CCACGCGAAC TCCGACCAGG ACGACCACGA CTTTCGCGACG GCCAGAAGCG  
 4951 CCTGCGCGTC GGCCAGGTAG CATTTGACCA TGGTGTGATA GTCCAGCCCC  
 GGACGCGCAG CCGGTCCATC GTAAACTGGT ACCACAGTAT CAGGTCGGGG  
 5001 TCCGCGGCGT GGCCCTTGCG GCGCAGCTTG CCCTTGAGAG AGGCGCCGCA  
 AGGCGCCGCA CCGGGAACCG CGCGTCGAAC GGGAACCTCC TCCGCGGCGT

FIG.9A-6

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5051 CGAGGGGCAG TGCAGACTTT TGAGGGCGTA GAGCTTGGGC GCGAGAAATA  
 GCTCCCCGTC ACGTCTGAAA ACTCCCGCAT CTCGAACCCG CGCTCTTTAT  
 5101 CCGATTCCGG GGAGTAGGCA TCCGCGCCGC AGGCCCCGCA GACGGTCTCG  
 GGCTAAGGCC CCTCATCCGT AGGCGCGGCG TCCGGGGCGT CTGCCAGAGC  
 5151 CATTCCACGA GCCAGGTGAG CTCTGGCCGT TCGGGGTCAA AAACCAGGTT  
 GTAAGGTGCT CGGTCCACTC GAGACCGGCA AGCCCCAGTT TTTGGTCCAA  
 5201 TCCCCCATGC TTTTGTATGC GTTTCTTACC TCTGGTTTCC ATGAGCCGGT  
 AGGGGGTACG AAAAAGTACG CAAAGAATGG AGACCAAAGG TACTCGGCCA  
 5251 GTCCACGCTC GGTGACGAAA AGGCTGTCCG TGTCCCCGTA TACAGACTTG  
 CAGGTGCGAG CCACTGCTTT TCCGACAGGC ACAGGGGCAT ATGTCTGAAC  
 5301 AGAGGCCTGT CCTCGAGCGG TGTTCCGCGG TCCTCCTCGT ATAGAAACTC  
 TCTCCGGACA GGAGCTCGCC ACAAGGCGCC AGGAGGAGCA TATCTTTGAG  
 5351 GGACCACTCT GAGACAAAGG CTCGCGTCCA GGCCAGCACG AAGGAGGCTA  
 CCTGGTGAGA CTCTGTTTCC GAGCGCAGGT CCGGTCGTGC TTCCTCCGAT  
 5401 AGTGGGAGGG GTAGCGGTCTG TTGTCCACTA GGGGGTCCAC TCGCTCCAGG  
 TCACCCTCCC CATCGCCAGC AACAGGTGAT CCCCAGGTG AGCGAGGTCC  
 5451 GTGTGAAGAC ACATGTCGCC CTCTTCGGCA TCAAGGAAGG TGATTGGTTT  
 CACACTTCTG TGTACAGCGG GAGAAGCCGT AGTTCCTTCC ACTAACCAA  
 5501 GTAGGTGTAG GCCACGTGAC CGGGTGTTCC TGAAGGGGGG CTATAAAAGG  
 CATCCACATC CGGTGCACTG GCCACAAGG ACTTCCCCC GATATTTTCC  
 5551 GGGTGGGGGC GCGTTCGTCC TCACTCTCTT CCGCATCGCT GTCTGCGAGG  
 CCCACCCCGG CGCAAGCAGG AGTGAGAGAA GGCGTAGCGA CAGACGCTCC  
 5601 GCCAGCTGTT GGGGTGAGTA CTCCCTCTGA AAAGCGGGCA TGACTTCTGC  
 CGGTCGACAA CCCCCTCAT GAGGGAGACT TTTCGCCCCT ACTGAAGACG  
 5651 GCTAAGATTG TCAGTTTCCA AAAACGAGGA GGATTTGATA TTCACCTGGC  
 CGATTCTAAC AGTCAAAGGT TTTTGCTCCT CCTAAACTAT AAGTGGACCG  
 5701 CCGCGGTGAT GCCTTTGAGG GTGGCCGCAT CCATCTGGTC AGAAAAGACA  
 GGCGCCACTA CGGAAACTCC CACCGGCGTA GGTAGACCAG TCTTTTCTGT  
 5751 ATCTTTTTGT TGTCAAGCTT GGTGGCAAAC GACCCGTAGA GGGCGTTGGA  
 TAGAAAAACA ACAGTTCGAA CCACCGTTTG CTGGGCATCT CCCGCAACCT  
 5801 CAGCAACTTG GCGATGGAGC GCAGGGTTTG GTTTTTGTCT CGATCGGCGC  
 GTCGTTGAAC CGCTACCTCG CGTCCCAAAC CAAAAACAGC GCTAGCCGCG  
 5851 GCTCCTTGGC CGCGATGTTT AGCTGCACGT ATTCGCGCGC AACGCACCGC  
 CGAGGAACCG GCGCTACAAA TCGACGTGCA TAAGCGCGCG TTGCGTGGCG

FIG.9A-7

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|      |            |             |             |             |             |
|------|------------|-------------|-------------|-------------|-------------|
| 5901 | CATTCGGGAA | AGACGGTGGT  | GCGCTCGTCG  | GGCACCAGGT  | GCACGCGCCA  |
|      | GTAAGCCCTT | TCTGCCACCA  | CGCGAGCAGC  | CCGTGGTCCA  | CGTGCGCGGT  |
| 5951 | ACCGCGGTTG | TGCAGGGTGA  | CAAGGTCAAC  | GCTGGTGGCT  | ACCTCTCCGC  |
|      | TGGCGCCAAC | ACGTCCCACT  | GTTCCAGTTG  | CGACCACCGA  | TGGAGAGGCG  |
| 6001 | GTAGGCGCTC | GTTGGTCCAG  | CAGAGGCGGC  | CGCCCTTGCG  | CGAGCAGAAAT |
|      | CATCCGCGAG | CAACCAGGTC  | GTCTCCGCCG  | GCGGGAACGC  | GCTCGTCTTA  |
| 6051 | GGCGGTAGGG | GGTCTAGCTG  | CGTCTCGTCC  | GGGGGGTCTG  | CGTCCACGGT  |
|      | CCGCCATCCC | CCAGATCGAC  | GCAGAGCAGG  | CCCCCAGAC   | GCAGGTGCCA  |
| 6101 | AAAGACCCCG | GGCAGCAGGC  | GCGCGTCGAA  | GTAGTCTATC  | TTGCATCCTT  |
|      | TTTCTGGGGC | CCGTCGTCCG  | CGCGCAGCTT  | CATCAGATAG  | AACGTAGGAA  |
| 6151 | GCAAGTCTAG | CGCCTGCTGC  | CATGCGCGGG  | CGGCAAGCGC  | GCGCTCGTAT  |
|      | CGTTCAGATC | GCGGACGACG  | GTACGCGCCC  | GCCGTTTCGCG | CGCGAGCATA  |
| 6201 | GGGTTGAGTG | GGGGACCCCA  | TGGCATGGGG  | TGGGTGAGCG  | CGGAGGCGTA  |
|      | CCCAACTCAC | CCCCTGGGGT  | ACCGTACCCC  | ACCCACTCGC  | GCCTCCGCAT  |
| 6251 | CATGCCGCAA | ATGTCGTAAA  | CGTAGAGGGG  | CTCTCTGAGT  | ATTCCAAGAT  |
|      | GTACGGCGTT | TACAGCATTT  | GCATCTCCCC  | GAGAGACTCA  | TAAGGTTCTA  |
| 6301 | ATGTAGGGTA | GCATCTTCCA  | CCGCGGATGC  | TGGCGCGCAC  | GTAATCGTAT  |
|      | TACATCCCAT | CGTAGAAGGT  | GGCGCCTACG  | ACCGCGCGTG  | CATTAGCATA  |
| 6351 | AGTTCGTGCG | AGGGAGCGAG  | GAGGTCGGGA  | CCGAGGTTGC  | TACGGGCGGG  |
|      | TCAAGCACGC | TCCCTCGCTC  | CTCCAGCCCT  | GGCTCCAACG  | ATGCCCGCCC  |
| 6401 | CTGCTCTGCT | CGGAAGACTA  | TCTGCCTGAA  | GATGGCATGT  | GAGTTGGATG  |
|      | GACGAGACGA | GCCTTCTGAT  | AGACGGACTT  | CTACCGTACA  | CTCAACCTAC  |
| 6451 | ATATGGTTGG | ACGCTGGAAG  | ACGTTGAAGC  | TGGCGTCTGT  | GAGACCTACC  |
|      | TATACCAACC | TGCGACCTTC  | TGCAACTTCG  | ACCGCAGACA  | CTCTGGATGG  |
| 6501 | GCGTCACGCA | CGAAGGAGGC  | GTAGGAGTCG  | CGCAGCTTGT  | TGACCAGCTC  |
|      | CGCAGTGCGT | GCTTCCTCCG  | CATCCTCAGC  | GCGTCGAACA  | ACTGGTCGAG  |
| 6551 | GGCGGTGACC | TGCACGTCTA  | GGGCGCAGTA  | GTCCAGGGTT  | TCCTTGATGA  |
|      | CCGCCACTGG | ACGTGCAGAT  | CCGCGTCAT   | CAGGTCCCAA  | AGGAACTACT  |
| 6601 | TGTCATACTT | ATCCTGTCCC  | TTTTTTTTTCC | ACAGCTCGCG  | GTTGAGGACA  |
|      | ACAGTATGAA | TAGGACAGGG  | AAAAAAAAGG  | TGTCGAGCGC  | CAACTCCTGT  |
| 6651 | AACTCTTCGC | GGTCTTTCCA  | GTA CTCTTGG | ATCGGAAACC  | CGTCGGCCTC  |
|      | TTGAGAAGCG | CCAGAAAAGGT | CATGAGAACC  | TAGCCTTTGG  | GCAGCCGGAG  |
| 6701 | CGAACGGTAA | GAGCCTAGCA  | TGTAGAACTG  | GTTGACGGCC  | TGGTAGGCGC  |
|      | GCTTGCCATT | CTCGGATCGT  | ACATCTTGAC  | CAACTGCCGG  | ACCATCCGCG  |

FIG.9A-8



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6751 AGCATCCCTT TTCTACGGGT AGCGCGTATG CCTGCGCGGC CTTCCGGAGC  
 TCGTAGGGAA AAGATGCCCA TCGCGCATAC GGACGCGCCG GAAGGCCTCG  
 6801 GAGGTGTGGG TGAGCGCAAA GGTGTCCCTG ACCATGACTT TGAGGTACTG  
 CTCCACACCC ACTCGCGTTT CCACAGGGAC TGGTACTGAA ACTCCATGAC  
 6851 GTATTTGAAG TCAGTGTCGT CGCATCCGCC CTGCTCCCAG AGCAAAAAGT  
 CATAAACTTC AGTCACAGCA GCGTAGGCGG GACGAGGGTC TCGTTTTTCA  
 6901 CCGTGCGCTT TTTGGAACGC GGATTTGGCA GGGCGAAGGT GACATCGTTG  
 GGCACGCGAA AAACCTTGCG CCTAAACCGT CCCGCTTCCA CTGTAGCAAC  
 6951 AAGAGTATCT TTCCGCGCG AGGCATAAAG TTGCGTGTGA TGCGBAAGGG  
 TTCTCATAGA AAGGGCGCGC TCCGTATTTC AACGCACACT ACGCCTTCCC  
 7001 TCCCGGCACC TCGGAACGGT TGTTAATTAC CTGGGCGGCG AGCACGATCT  
 AGGGCCGTGG AGCCTTGCCA ACAATTAATG GACCCGCCGC TCGTGCTAGA  
 7051 CGTCAAAGCC GTTGATGTTG TGGCCACAA TGTAAGTTC CAAGAAGCGC  
 GCAGTTTCGG CAACTACAAC ACCGGGTGTT ACATTTCAAG GTTCTTCGCG  
 7101 GGGATGCCCT TGATGGAAGG CAATTTTTTA AGTTCCTCGT AGGTGAGCTC  
 CCCTACGGGA ACTACCTTCC GTTAAAAAAT TCAAGGAGCA TCCACTCGAG  
 7151 TTCAGGGGAG CTGAGCCCGT GCTCTGAAAG GGCCAGTCT GCAAGATGAG  
 AAGTCCCCTC GACTCGGGCA CGAGACTTTC CCGGGTCAGA CGTTCTACTC  
 7201 GGTTGGAAGC GACGAATGAG CTCCACAGGT CACGGGCCAT TAGCATTTGC  
 CCAACCTTCG CTGCTTACTC GAGGTGTCCA GTGCCCGGTA ATCGTAAACG  
 7251 AGGTGGTCGC GAAAGGTCCT AAAGTGGCGA CCTATGGCCA TTTTTTCTGG  
 TCCACCAGCG CTTTCCAGGA TTTGACCGCT GGATACCGGT AAAAAAGACC  
 7301 GGTGATGCAG TAGAAGGTAA GCGGGTCTTG TTCCAGCGG TCCCATCCAA  
 CCACTACGTC ATCTTCCATT CGCCCAGAAC AAGGGTCGCC AGGGTAGGTT  
 7351 GGTTGCGGCG TAGGTCTCGC GCGGCAGTCA CTAGAGGCTC ATCTCCGCCG  
 CCAAGCGCCG ATCCAGAGCG CGCCGTCACT GATCTCCGAG TAGAGGCGGC  
 7401 AACTTCATGA CCAGCATGAA GGGCACGAGC TGCTTCCCAA AGGCCCCCAT  
 TTGAAGTACT GGTCGTACTT CCCGTGCTCG ACGAAGGGTT TCCGGGGGTA  
 7451 CCAAGTATAG GTCTCTACAT CGTAGGTGAC AAAGAGACGC TCGGTGCGAG  
 GGTTTCATATC CAGAGATGTA GCATCCACTG TTTCTCTGCG AGCCACGCTC  
 7501 GATGCGAGCC GATCGGGAAG AACTGGATCT CCCGCCACCA ATTGGAGGAG  
 CTACGCTCGG CTAGCCCTTC TTGACCTAGA GGGCGGTGGT TAACCTCCTC  
 7551 TGGCTATTGA TGTGGTGAAA GTAGAAGTCC CTGCGACGGG CCGAACACTC  
 ACCGATAACT ACACCACTTT CATCTTCAGG GACGCTGCCG GGCTTGTGAG

FIG.9A-9

18/70

7601 GTGCTGGCTT TTGTAAAAAC GTGCGCAGTA CTGGCAGCGG TGCACGGGCT  
CACGACCGAA AACATTTTTG CACGCGTCAT GACCGTCGCC ACGTGCCCGA

7651 GTACATCCTG CACGAGGTTG ACCTGACGAC CGCGCACAAG GAAGCAGAGT  
CATGTAGGAC GTGCTCCAAC TGGACTGCTG GCGCGTGTTT CTTGCTCTCA

7701 GGGAATTTGA GCCCTCGCC TGGCGGGTTT GGCTGGTGGT CTTCTACTTC  
CCCTTAAACT CGGGGAGCGG ACCGCCAAA CCGACCACCA GAAGATGAAG

7751 GGCTGCTTGT CTTGACCGT CTGGCTGCTC GAGGGGAGTT ACGGTGGATC  
CCGACGAACA GGAAGTGGCA GACCGACGAG CTCCCTCAA TGCCACCTAG

7801 GGACCACCAC GCCGCGCGAG CCCAAAGTCC AGATGTCCGC GCGCGGCGGT  
CCTGGTGGTG CGGCGCGCTC GGGTTTCAGG TCTACAGGCG CGCGCCGCCA

7851 CGGAGCTTGA TGACAACATC GCGCAGATGG GAGCTGTCCA TGGTCTGGAG  
GCCTCGAACT ACTGTTGTAG CGCGTCTACC CTCGACAGGT ACCAGACCTC

7901 CTCCCGCGGC GTCAGGTCAG GCGGGAGCTC CTGCAGGTTT ACCTCGCATA  
GAGGGCGCCG CAGTCCAGTC CGCCCTCGAG GACGTCCAAA TGGAGCGTAT

7951 GACGGGTCAG GCGCGGGGCT AGATCCAGGT GATACCTAAT TTCCAGGGGC  
CTGCCAGTC CCGCGCCCGA TCTAGGTCCA CTATGGATTA AAGGTCCCCG

8001 TGGTTGGTGG CGGCGTCGAT GGCTTGCAAG AGGCCGCATC CCCGCGGCGC  
ACCAACCACC GCCGCAGCTA CCGAACGTTT TCCGGCGTAG GGGCGCCGCG

8051 GACTACGGTA CCGCGCGGCG GCGGGTGGGC CGCGGGGGTG TCCTTGGATG  
CTGATGCCAT GCGCGCGCCG CCGCCACCCG GCGCCCCAC AGGAACCTAC

8101 ATGCATCTAA AAGCGGTGAC GCGGGCGAGC CCGCGGAGGT AGGGGGGGCT  
TACGTAGATT TTCGCCACTG CGCCCGCTCG GGGGCTCCA TCCCCCGA

8151 CCGGACCCGC CGGGAGAGGG GGCAGGGGCA CGTCGGCGCC GCGCGCGGGC  
GGCCTGGGCG GCCCTCTCC CCGTCCCCGT GCAGCCGCGG CGCGCGCCCG

8201 AGGAGCTGGT GCTGCGCGCG TAGGTTGCTG GCGAACGCGA CGACGCGGCG  
TCCTCGACCA CGACGCGCGC ATCCAACGAC CGCTTGCGCT GCTGCGCCGC

8251 GTTGATCTCC TGAATCTGGC GCCTCTGCGT GAAGACGACG GGCCCGGTGA  
CAACTAGAGG ACTTAGACCG CGGAGACGCA CTTCTGCTGC CCGGGCCACT

8301 GCTTGAACCT GAAAGAGAGT TCGACAGAAT CAATTTGCGT GTCGTTGACG  
CGAACTTGA CTTTCTCTCA AGCTGTCTTA GTTAAAGCCA CAGCAACTGC

8351 GCGGCCTGGC GCAAAATCTC CTGCACGTCT CCTGAGTTGT CTTGATAGGC  
CGCCGGACCG CGTTTTAGAG GACGTGCAGA GGAACAACA GAAGTATCCG

8401 GATCTCGGCC ATGAACTGCT CGATCTCTTC CTCCTGGAGA TCTCCGCGTC  
CTAGAGCCGG TACTTGACGA GCTAGAGAAG GAGGACCTCT AGAGGCGCAG

FIG.9A-10

19/70

8451 CGGCTCGCTC CACGGTGGCG GCGAGGTCGT TGGAAATGCG GGCCATGAGC  
GCCGAGCGAG GTGCCACCGC CGCTCCAGCA ACCTTTACGC CCGGTACTCG

8501 TGCAGAGAAGG CGTTGAGGCC TCCCTCGTTC CAGACGCGGC TGTAGACCAC  
ACGCTCTTCC GCAACTCCGG AGGGAGCAAG GTCTGCGCCG ACATCTGGTG

8551 GCCCCCTTCG GCATCGCGGG CGCGCATGAC CACCTGCGCG AGATTGAGCT  
CGGGGAAGC CGTAGCGCCC GCGCGTACTG GTGGACGCGC TCTAACTCGA

8601 CCACGTGCCG GGCAGAGACG GCGTAGTTTC GCAGGCGCTG AAAGAGGTAG  
GGTGACGGC CCGCTTCTGC CGCATCAAAG CGTCCGCGAC TTTCTCCATC

8651 TTGAGGGTGG TGGCGGTGTG TTCTGCCACG AAGAAGTACA TAACCCAGCG  
AACTCCACC ACCGCCACAC AAGACGGTGC TTCTTCATGT ATTGGGTCGC

8701 TCGCAACGTG GATTCGTTGA TATCCCCCAA GGCCTCAAGG CGCTCCATGG  
AGCGTTGCAC CTAAGCAACT ATAGGGGGTT CCGGAGTTCC GCGAGGTACC

8751 CCTCGTAGAA GTCCACGGCG AAGTTGAAAA ACTGGGAGTT GCGCGCCGAC  
GGAGCATCTT CAGGTGCCGC TTCAACTTTT TGACCCTCAA CGCGCGGCTG

8801 ACGGTAACT CCTCCTCCAG AAGACGGATG AGCTCGGCGA CAGTGTGCGG  
TGCCAATTGA GGAGGAGGTC TTCTGCCTAC TCGAGCCGCT GTCACAGCGC

8851 CACCTCGCGC TCAAAGGCTA CAGGGGCCTC TTCTTCTTCT TCAATCTCCT  
GTGGAGCGCG AGTTTCCGAT GTCCCCGGAG AAGAAGAAGA AGTTAGAGGA

8901 CTTCCATAAG GGCTCCCCT TCTTCTTCTT CTGGCGGCGG TGGGGGAGGG  
GAAGGTATTG CCGGAGGGGA AGAAGAAGAA GACCGCCGCC ACCCCCTCCC

8951 GGGACACGGC GCGCAGCAGC GCGCACCAGG AGGCGGTGCA CAAAGCGCTC  
CCCTGTGCCG CCGCTGCTGC CGCGTGGCCC TCCGCCAGCT GTTTCGCGAG

9001 GATCATCTCC CCGCGGCGAC GCGCATGGT CTCGGTGACG GCGCGGCCGT  
CTAGTAGAGG GCGCCGCTG CCGGTACCA GAGCCACTGC CGCGCCGGCA

9051 TCTCGCGGGG GCGCAGTTGG AAGACGCCGC CCGTCATGTC CCGGTTATGG  
AGAGCGCCCC CGCGTCAACC TTCTGCGGCG GGCAGTACAG GGCCAATACC

9101 GTTGGCGGGG GGCTGCCATG CGGCAGGGAT ACGGCGCTAA CGATGCATCT  
CAACCGCCCC CCGACGGTAC GCCGTCCCTA TGCCGCGATT GCTACGTAGA

9151 CAACAATTGT TGTGTAGGTA CTCCGCCGCC GAGGGACCTG AGCGAGTCCG  
GTTGTTAACA ACACATCCAT GAGGCGGCGG CTCCCTGGAC TCGCTCAGGC

9201 CATCGACCGG ATCGGAAAAC CTCTCGAGAA AGGCGTCTAA CCAGTCACAG  
GTAGCTGGCC TAGCCTTTTG GAGAGCTCTT TCCGAGATT GGTCAGTGTC

9251 TCGCAAGGTA GGCTGAGCAC CGTGGCGGGC GGCAGCGGGC GGCAGTCGGG  
AGCGTTCCAT CCGACTCGTG GCACCGCCCG CCGTCGCCCC CCGCCAGCCC

FIG.9A-11

20/70

9301 GTTGTCTTCTG GCGGAGGTGC TGCTGATGAT GTAATTAAAG TAGGCGGTCT  
 CAACAAAGAC CGCCTCCACG ACGACTACTA CATTAAATTC ATCCGCCAGA  
 9351 TGAGACGGCG GATGGTCGAC AGAAGCACCA TGTCTTGGG TCCGGCCTGC  
 ACTCTGCCGC CTACCAGCTG TCTTCGTGGT ACAGGAACCC AGGCCGGACG  
 9401 TGAATGCGCA GGC GGTCGGC CATGCCCCAG GCTTCGTTTT GACATCGGCG  
 ACTTACGCGT CCGCCAGCCG GTACGGGGTC CGAAGCAAAA CTGTAGCCGC  
 9451 CAGGTCTTTG TAGTAGTCTT GCATGAGCCT TTCTACCGGC ACTTCTTCTT  
 GTCCAGAAAC ATCATCAGAA CGTACTCGGA AAGATGGCCG TGAAGAAGAA  
 9501 CTCCTTCCTC TTGTCCTGCA TCTCTTGCAT CTATCGCTGC GGCGGCGGCG  
 GAGGAAGGAG AACAGGACGT AGAGAACGTA GATAGCGACG CCGCCGCCGC  
 9551 GAGTTTGGCC GTAGGTGGCG CCCTCTTCCT CCCATGCGTG TGACCCCGAA  
 CTCAAACCGG CATCCACCGC GGGAGAAGGA GGGTACGCAC ACTGGGGCTT  
 9601 GCCCCTCATC GGCTGAAGCA GGGCTAGGTC GGCGACAACG CGCTCGGCTA  
 CGGGGAGTAG CCGACTTCGT CCCGATCCAG CCGCTGTTGC GCGAGCCGAT  
 9651 ATATGGCCTG CTGCACCTGC GTGAGGGTAG ACTGGAAGTC ATCCATGTCC  
 TATACCGGAC GACGTGGACG CACTCCCATC TGACCTTCAG TAGGTACAGG  
 9701 ACAAAGCGGT GGTATGCGCC CGTGTTGATG GTGTAAGTGC AGTTGGCCAT  
 TGTTTCGCCA CCATACGCGG GCACAACTAC CACATTCACG TCAACCGGTA  
 9751 AACGGACCAG TTAACGGTCT GGTGACCCGG CTGCGAGAGC TCGGTGTACC  
 TTGCCTGGTC AATTGCCAGA CCACTGGGCC GACGCTCTCG AGCCACATGG  
 9801 TGAGACGCGA GTAAGCCCTC GAGTCAAATA CGTAGTCGTT GCAAGTCCGC  
 ACTCTGCGCT CATTCGGGAG CTCAGTTTAT GCATCAGCAA CGTTCAGGCG  
 9851 ACCAGGTACT GGTATCCCAC CAAAAAGTGC GGCGGCGGCT GGCGGTAGAG  
 TGGTCCATGA CCATAGGGTG GTTTTTACG CCGCCGCCGA CCGCCATCTC  
 9901 GGGCCAGCGT AGGGTGGCCG GGGCTCCGGG GGCGAGATCT TCCAACATAA  
 CCCGGTCGCA TCCCACCGGC CCCGAGGCC CCGCTCTAGA AGGTTGTATT  
 9951 GGCGATGATA TCCGTAGATG TACCTGGACA TCCAGGTGAT GCCGGCGGCG  
 CCGCTACTAT AGGCATCTAC ATGGACCTGT AGGTCCACTA CGGCCGCCGC  
 10001 GTGGTGGAGG CGCGCGGAAA GTCGCGGACG CGGTTCCAGA TGTTGCGCAG  
 CACCACCTCC GCGCGCCTTT CAGCGCCTGC GCCAAGGTCT ACAACGCGTC  
 10051 CGGCAAAAAG TGCTCCATGG TCGGGACGCT CTGGCCGGTC AGGCGCGCGC  
 GCCGTTTTTC ACGAGGTACC AGCCCTGCGA GACCGGCCAG TCCGCGCGCG  
 10101 AATCGTTGAC GCTCTAGACC GTGCAAAAGG AGAGCCTGTA AGCGGGCACT  
 TTAGCAACTG CGAGATCTGG CACGTTTTCC TCTCGGACAT TCGCCCGTGA

FIG.9A-12

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10151 CTTCCGTGGT CTGGTGGATA AATTCGCAAG GGTATCATGG CGGACGACCG  
 GAAGGCACCA GACCACCTAT TTAAGCGTTC CCATAGTACC GCCTGCTGGC  
 10201 GGGTTCGAGC CCCGTATCCG GCCGTCCGCC GTGATCCATG CGGTTACCGC  
 CCCAAGCTCG GGGCATAGGC CGGCAGGCGG CACTAGGTAC GCCAATGGCG  
 10251 CCGCGTGTCG AACCCAGGTG TGCACGTCA GACAACGGGG GAGTGCTCCT  
 GGCACACAGC TTGGGTCCAC ACGCTGCACT CTGTTGCCCC CTCACGAGGA  
 10301 TTTGGCTTCC TTCCAGGCGC GCGGCTGCT GCGCTAGCTT TTTTGGCCAC  
 AAACCGAAGG AAGGTCCGCG CCGCCGACGA CCGATCGAA AAAACCGGTG  
 10351 TGGCCGCGCG CAGCGTAAGC GGTTAGGCTG GAAAGCGAAA GCATTAAGTG  
 ACCGGCGCGC GTCGCATTG CCAATCCGAC CTTTCGCTTT CGTAATTAC  
 10401 GCTCGCTCCC TGTAGCCGGA GGGTTATTTT CCAAGGGTTG AGTCGCGGGA  
 CGAGCGAGGG ACATCGGCCCT CCAATAAAA GGTTCCTAAC TCAGCGCCCT  
 10451 CCCCCGGTTC GAGTCTCGGA CCGGCCGGAC TCGGGCGAAC GGGGGTTTGC  
 GGGGGCCAAG CTCAGAGCCT GGCCGGCCTG ACGCCGCTTG CCCCCAAACG  
 10501 CTCCCCGTCA TGCAAGACCC CGCTTGCAAA TTCCTCCGGA AACAGGGACG  
 GAGGGGCAGT ACGTTCTGGG GCGAACGTTT AAGGAGGCCT TTGTCCCTGC  
 10551 AGCCCCTTTT TTGCTTTTCC CAGATGCATC CCGTGCTGCG GCAGATGCGC  
 TCGGGGAAAA AACGAAAAGG GTCTACGTAG GCCACGACGC CGTCTACGCG  
 10601 CCCCCTCCTC AGCAGCGGCA AGAGCAAGAG CAGCGGCAGA CATGCAGGGC  
 GGGGGAGGAG TCGTCGCCGT TCTCGTTCTC GTCGCCGTCT GTACGTCCCG  
 10651 ACCCTCCCCT CCTCCTACCG CGTCAGGAGG GGCACATCC GCGGTTGACG  
 TGGGAGGGGA GGAGGATGGC GCAGTCCTCC CCGCTGTAGG CGCCAAGTGC  
 10701 CCGCAGCAGA TGGTGATTAC GAACCCCCGC GCGGCCGGGC CCGGCACTAC  
 GCCGTGCTCT ACCACTAATG CTTGGGGGCG CCGCGGCCCG GGCCGTGATG  
 10751 CTGGACTTGG AGGAGGGCGA GGGCCTGGCG CCGCTAGGAG CGCCCTCTCC  
 GACCTGAACC TCCTCCCGCT CCCGACCGC GCCGATCCTC GCGGGAGAGG  
 10801 TGAGCGGCAC CCAAGGGTGC AGCTGAAGCG TGATACGCGT GAGGCGTACG  
 ACTCGCCGTG GGTTCCACG TCGACTTCGC ACTATGCGCA CTCCGCATGC  
 10851 TGCCGCGGCA GAACCTGTTT CCGCACCAGC AGGGAGAGGA GCCCGAGGAG  
 ACGGCGCCGT CTTGGACAAA GCGCTGGCGC TCCCTCTCCT CGGGCTCCTC  
 10901 ATGCGGGATC GAAAGTTCCA CGCAGGGCGC GAGCTGCGGC ATGGCCTGAA  
 TACGCCCTAG CTTTCAAGGT GCGTCCCGCG CTCGACGCCG TACCGGACTT  
 10951 TCGCGAGCGG TTGCTGCGCG AGGAGGACTT TGAGCCCGAC GCGCGAACC  
 AGCGCTCGCC AACGACGCGC TCCTCCTGAA ACTCGGGCTG CCGCTTGGC

FIG.9A-13

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11001 GGATTAGTCC CGCGCGCGCA CACGTGGCGG CCGCCGACCT GGTAACCGCA  
 CCTAATCAGG GCGCGCGCGT GTGCACCGCC GCGGGCTGGA CCATTGGCGT

11051 TACGAGCAGA CGGTGAACCA GGAGATTAAC TTTCAAAAAA GCTTTAACAA  
 ATGCTCGTCT GCCACTTGGT CCTCTAATTG AAAGTTTTTT CGAAATTGTT

11101 CCACGTGCGT ACGCTTGTGG CGCGCGAGGA GGTGGCTATA GGA CTGATGC  
 GGTGCACGCA TCGGAACACC GCGCGCTCCT CCACCGATAT CCTGACTACG

11151 ATCTGTGGGA CTTTGTAAGC GCGCTGGAGC AAAACCCAAA TAGCAAGCCG  
 TAGACACCTT GAAACATTG GCGGACCTCG TTTTGGGTTT ATCGTTCGGC

11201 CTCATGGCGC AGCTGTTCTT TATAGTGCAG CACAGCAGGG ACAACGAGGC  
 GAGTACCGCG TCGACAAGGA ATATCACGTC GTGTCGTCCT TGTGCTCCG

11251 ATTCAGGGAT GCGCTGCTAA ACATAGTAGA GCGCGAGGGC CGCTGGCTGC  
 TAAGTCCCTA CGCGACGATT TGTATCATCT CCGGCTCCCG GCGACCGACG

11301 TCGATTTGAT AAACATCCTG CAGAGCATAG TGGTGCAGGA GCGCAGCTTG  
 AGCTAAACTA TTTGTAGGAC GTCTCGTATC ACCACGTCCT CGCGTCGAAC

11351 AGCCTGGCTG ACAAGGTGGC CGCCATCAAC TATTCCATGC TTAGCCTGGG  
 TCGGACCGAC TGTTCCACCG GCGGTAGTTG ATAAGGTACG AATCGGACCC

11401 CAAGTTTTAC GCGCGCAAGA TATACCATAC CCCTTACGTT CCCATAGACA  
 GTTCAAAATG CGGGCGTTCT ATATGGTATG GGAATGCAA GGGTATCTGT

11451 AGGAGGTAAA GATCGAGGGG TTCTACATGC GCATGGCGCT GAAGGTGCTT  
 TCCTCCATTT CTAGTCCCCC AAGATGTACG CGTACCGCGA CTTCCACGAA

11501 ACCTTGAGCG ACGACCTGGG CGTTTATCGC AACGAGCGCA TCCACAAGGC  
 TGGAACTCGC TGCTGGACCC GCAAATAGCG TTGCTCGCGT AGGTGTTCCG

11551 CGTGAGCGTG AGCCGGCGGC GCGAGCTCAG CGACCGCGAG CTGATGCACA  
 GCACTCGCAC TCGGCCGCCG CGCTCGAGTC GCTGGCGCTC GACTACGTGT

11601 GCCTGCAAAG GGCCCTGGCT GGCACGGGCA GCGGCGATAG AGAGGCCGAG  
 CGGACGTTTC CCGGGACCGA CCGTGCCCGT CGCCGCTATC TCTCCGGCTC

11651 TCCTACTTTG ACGCGGGCGC TGACCTGCGC TGGGCCCCAA GCCGACGCGC  
 AGGATGAAAC TGCGCCCGCG ACTGGACGCG ACCCGGGGTT CGGCTGCGCG

11701 CCTGGAGGCA GCTGGGGCCG GACCTGGGCT GGCAGTGGCA CCCGCGCGCG  
 GGACCTCCGT CGACCCCGGC CTGGACCCGA CCGCCACCGT GGGCGCGCGC

11751 CTGGCAACGT CGGCGGCGTG GAGGAATATG ACGAGGACGA TGAGTACGAG  
 GACCGTTGCA GCCGCCGCAC CTCCTTATAC TGCTCCTGCT ACTCATGCTC

11801 CCAGAGGACG GCGAGTACTA AGCGGTGATG TTTCTGATCA GATGATGCAA  
 GGTCTCCTGC CGCTCATGAT TCGCCACTAC AAAGACTAGT CTACTACGTT

FIG.9A-14

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11851 GACGCAACGG ACCCGGCGGT GCGGGCGGCG CTGCAGAGCC AGCCGTCCGG  
 CTGCGTTGCC TGGGCCGCCA CGCCGCGCG GACGTCTCGG TCGGCAGGCC  
 11901 CCTTAACTCC ACGGACGACT GGC GCCAGGT CATGGACCGC ATCATGTGCG  
 GGAATTGAGG TGCCTGCTGA CCGCGGTCCA GTACCTGGCG TAGTACAGCG  
 11951 TGA CTGCGCG CAATCCTGAC GCGTTCCGGC AGCAGCCGCA GGCCAACCGG  
 ACTGACGCGC GTTAGGACTG CGCAAGGCCG TCGTCGGCGT CCGGTTGGCC  
 12001 CTCTCCGCAA TTCTGGAAGC GGTGGTCCCG GCGCGCGCAA ACCCCACGCA  
 GAGAGGCGTT AAGACCTTCG CCACCAGGGC CCGCGCGGTT TGGGGTGCGT  
 12051 CGAGAAGGTG CTGGCGATCG TAAACGCGCT GGCCGAAAAC AGGGCCATCC  
 GCTCTTCCAC GACCGCTAGC ATTTGCGCGA CCGGCTTTTG TCCCGGTAGG  
 12101 GGCCCGACGA GGCCGGCCTG GTCTACGACG CGCTGCTTCA GCGCGTGGCT  
 CCGGGCTGCT CCGGCCGGAC CAGATGCTGC GCGACGAAGT CGCGCACCGA  
 12151 CGTTACAACA GCGGCAACGT GCAGACCAAC CTGGACCGGC TGGTGGGGGA  
 GCAATGTTGT CGCCGTTGCA CGTCTGGTTG GACCTGGCCG ACCACCCCT  
 12201 TGTGCGCGAG GCCGTGGCGC AGCGTGAGCG CGCGCAGCAG CAGGGCAACC  
 ACACGCGCTC CGGCACCGCG TCGCACTCGC GCGCGTCGTC GTCCCGTTGG  
 12251 TGGGCTCCAT GGTGCTACTA AACGCCTTCC TGAGTACACA GCCCGCCAAC  
 ACCCGAGGTA CCAACGTGAT TTGCGGAAGG ACTCATGTGT CGGGCGGTTG  
 12301 GTGCCGCGGG GACAGGAGGA CTACACCAAC TTTGTGAGCG CACTGCGGCT  
 CACGGCGCCC CTGTCTCCT GATGTGGTTG AAACACTCGC GTGACGCCGA  
 12351 AATGGTGACT GAGACACCGC AAAGTGAGGT GTACCACTCT GGGCCAGACT  
 TTACCACTGA CTCTGTGGCG TTTCACTCCA CATGGTCAGA CCCGGTCTGA  
 12401 ATTTTTTCCA GACCAGTAGA CAAGGCCTGC AGACCGTAAA CCTGAGCCAG  
 TAAAAAAGGT CTGGTCATCT GTTCCGGACG TCTGGCATTG GACTCGGTC  
 12451 GCTTTCAAAA ACTTGACGGG GCTGTGGGGG GTGCGGGCTC CCACAGGCGA  
 CGAAAGTTTT TGAACGTCCC CGACACCCCC CACGCCCAG GGTGTCCGCT  
 12501 CCGCGCGACC GTGTCTAGCT TGCTGACGCC CAACTCGCGC CTGTTGCTGC  
 GGCGCGCTGG CACAGATCGA ACGACTGCGG GTTGAGCGCG GACAACGACG  
 12551 TGCTAATAGC GCCCTTCACG GACAGTGGCA GCGTGTCCCG GGACACATAC  
 ACGATTATCG CGGGAAGTGC CTGTCAACGT CGCACAGGGC CCTGTGTATG  
 12601 CTAGGTCCT TGCTGACACT GTACCGCGAG GCCATAGGTC AGGCGCATGT  
 GATCCAGTGA ACGACTGTGA CATGGCGCTC CCGTATCCAG TCCGCGTACA  
 12651 GGACGAGCAT ACTTTCAGG AGATTACAAG TGTCAGCCGC GCGCTGGGGC  
 CCTGCTCGTA TGAAAGGTCC TCTAATGTTT ACAGTCGGCG CGCGACCCCG

FIG.9A-15

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12701 AGGAGGACAC GGGCAGCCTG GAGGCAACCC TAAACTACCT GCTGACCAAC  
 TCCTCCTGTG CCCGTCGGAC CTCCGTTGGG ATTTGATGGA CGACTGGTTG  
 12751 CGGCGGCAGA AGATCCCCTC GTTGCACAGT TTAAACAGCG AGGAGGAGCG  
 GCCGCCGTCT TCTAGGGGAG CAACGTGTCA AATTTGTCGC TCCTCCTCGC  
 12801 CATTTTGCGC TACGTGCAGC AGAGCGTGAG CCTTAACCTG ATGCGCGACG  
 GTAAAACGCG ATGCACGTCG TCTCGCACTC GGAATTGGAC TACGCGCTGC  
 12851 GGGTAACGCC CAGCGTGGCG CTGGACATGA CCGCGCGCAA CATGGAACCG  
 CCCATTGCGG GTCGCACCGC GACCTGTACT GGC GCGCGTT GTACCTTGGC  
 12901 GGCATGTATG CCTCAAACCG GCCGTTTATC AACCGCCTAA TGGACTACTT  
 CCGTACATAC GGAGTTTGGC CGGCAAATAG TTGGCGGATT ACCTGATGAA  
 12951 GCATCGCGCG GCCGCCGTGA ACCCGAGTA TTTCACCAAT GCCATCTTGA  
 CGTAGCGCGC CGCGGGCACT TGGGGCTCAT AAAGTGGTTA CGGTAGAACT  
 13001 ACCCGCACTG GCTACCGCCC CCTGGTTTCT ACACCGGGGG ATTCGAGGTG  
 TGGGCGTGAC CGATGGCGGG GGACCAAAGA TGTGGCCCCC TAAGCTCCAC  
 13051 CCCGAGGGTA ACGATGGATT CCTCTGGGAC GACATAGACG ACAGCGTGTT  
 GGGCTCCCAT TGCTACCTAA GGAGACCCTG CTGTATCTGC TGTCGCACAA  
 13101 TTCCCCGCAA CCGCAGACCC TGCTAGAGTT GCAACAGCGC GAGCAGGCAG  
 AAGGGGCGTT GCGTCTGGG ACGATCTCAA CGTTGTCGCG CTCGTCCGTC  
 13151 AGGCGGCGCT GCGAAAGGAA AGCTTCCGCA GGCCAAGCAG CTTGTCCGAT  
 TCCGCCGCGA CGCTTTCCTT TCGAAGGCGT CCGGTTTCGC GAACAGGCTA  
 13201 CTAGGCGCTG CGGCCCGCG GTCAGATGCT AGTAGCCCAT TTCCAAGCTT  
 GATCCGCGAC GCCGGGGCGC CAGTCTACGA TCATCGGGTA AAGGTTTCGAA  
 13251 GATAGGGTCT CTTACCAGCA CTCGACCAC CCGCCCGCGC CTGCTGGGCG  
 CTATCCCAGA GAATGGTCGT GAGCGTGGTG GCGGGGCGCG GACGACCCGC  
 13301 AGGAGGAGTA CCTAAACAAC TCGTGCTGC AGCCGCAGCG CGAAAAAAAC  
 TCCTCCTCAT GGATTTGTTG AGCGACGACG TCGGCGTCGC GCTTTTTTTG  
 13351 CTGCCTCCGG CATTTCCCAA CAACGGGATA GAGAGCCTAG TGGACAAGAT  
 GACGGAGGCC GTAAAGGGTT GTTGCCCTAT CTCTCGGATC ACCTGTTCTA  
 13401 GAGTAGATGG AAGACGTACG CGCAGGAGCA CAGGGACGTG CCAGGCCCCG  
 CTCATCTACC TTCTGCATGC GCGTCTCTGT GTCCCTGCAC GGTCCGGGCG  
 13451 GCCCGCCAC CCGTCGTCAA AGGCACGACC GTCAGCGGGG TCTGGTGTGG  
 CGGGCGGGTG GGCAGCAGTT TCCGTGCTGG CAGTCGCCCC AGACCACACC  
 13501 GAGGACGATG ACTCGGCAGA CGACAGCAGC GTCCTGGATT TGGGAGGGAG  
 CTCCTGTAC TGAGCCGTCT GCTGTGCTCG CAGGACCTAA ACCCTCCCTC

FIG.9A-16



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13551 TGGCAACCCG TTTGCGCACC TTCGCCCCAG GCTGGGGAGA ATGTTTTTAAA  
 ACCGTTGGGC AAACGCGTGG AAGCGGGGTC CGACCCCTCT TACAAAATTT  
 13601 AAAAAAAAAA GCATGATGCA AAATAAAAAA CTCACCAAGG CCATGGCACC  
 TTTTTTTTTT CGTACTACGT TTTATTTTTT GAGTGGTTCC GGTACCGTGG  
 13651 GAGCGTTGGT TTTCTTGTAT TCCCCTTAGT ATGCGGCGCG CGGCGATGTA  
 CTCGCAACCA AAAGAACATA AGGGGAATCA TACGCCGCGC GCCGCTACAT  
 13701 TGAGGAAGGT CCTCCTCCCT CCTACGAGAG TGTGGTGAGC GCGGCGCCAG  
 ACTCCTTCCA GGAGGAGGGA GGATGCTCTC ACACCACTCG CGCCGCGGTC  
 13751 TGGCGGCGGC GCTGGGTTCT CCCTTCGATG CTCCCCTGGA CCCGCCGTTT  
 ACCGCCGCCG CGACCCAAGA GGAAGCTAC GAGGGGACCT GGGCGGCAAA  
 13801 GTGCCTCCGC GGTACCTGCG GCCTACCGGG GGGAGAAACA GCATCCGTTA  
 CACGGAGGCG CCATGGACGC CGGATGGCCC CCCTCTTTGT CGTAGGCAAT  
 13851 CTCTGAGTTG GCACCCCTAT TCGACACCAC CCGTGTGTAC CTGGTGGACA  
 GAGACTCAAC CGTGGGGATA AGCTGTGGTG GGCACACATG GACCACCTGT  
 13901 ACAAGTCAAC GGATGTGGCA TCCCTGAACT ACCAGAACGA CCACAGCAAC  
 TGTTCAGTTG CCTACACCGT AGGGACTTGA TGGTCTTGCT GGTGTCGTTG  
 13951 TTTCTGACCA CGGTCATTCA AAACAATGAC TACAGCCCGG GGGAGGCAAG  
 AAAGACTGGT GCCAGTAAGT TTTGTTACTG ATGTCGGGCC CCCTCCGTTT  
 14001 CACACAGACC ATCAATCTTG ACGACCGGTC GCACTGGGGC GGCGACCTGA  
 GTGTGTCTGG TAGTTAGAAC TGCTGGCCAG CGTGACCCCG CCGCTGGACT  
 14051 AAACCATCCT GCATACCAAC ATGCCAAATG TGAACGAGTT CATGTTTACC  
 TTTGGTAGGA CGTATGGTTG TACGGTTTAC ACTTGCTCAA GTACAAATGG  
 14101 AATAAGTTTA AGGCGCGGGT GATGGTGTCT CGCTTGCCTA CTAAGGACAA  
 TTATTCAAAT TCCGCGCCCA CTACCACAGC GCGAACGGAT GATTCTGTGT  
 14151 TCAGGTGGAG CTGAAATACG AGTGGGTGGA GTTCACGCTG CCCGAGGGCA  
 AGTCCACCTC GACTTTATGC TCACCCACCT CAAGTGCGAC GGGCTCCCGT  
 14201 ACTACTCCGA GACCATGACC ATAGACCTTA TGAACAACGC GATCGTGGAG  
 TGATGAGGCT CTGGTACTGG TATCTGGAAT ACTTGTGCG CTAGCACCTC  
 14251 CACTACTTGA AAGTGGGCAG ACAGAACGGG GTTCTGGAAA GCGACATCGG  
 GTGATGAACT TTCACCCGTC TGTCTTGCCC CAAGACCTTT CGCTGTAGCC  
 14301 GGTAAGTTT GACACCCGCA ACTTCAGACT GGGGTTTGAC CCCGTCACTG  
 CCATTTCAAA CTGTGGGCGT TGAAGTCTGA CCCCAACTG GGGCAGTGAC  
 14351 GTCTTGTCAT GCCTGGGGTA TATACAAACG AAGCCTTCCA TCCAGACATC  
 CAGAACAGTA CGGACCCCAT ATATGTTTGC TTCGGAAGGT AGGTCTGTAG

FIG.9A-17

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|       |                           |                          |                           |                          |                          |
|-------|---------------------------|--------------------------|---------------------------|--------------------------|--------------------------|
| 14401 | ATTTTGCTGC<br>TAAAACGACG  | CAGGATGCGG<br>GTCCTACGCC | GGTGGACTTC<br>CCACCTGAAG  | ACCCACAGCC<br>TGGGTGTCGG | GCCTGAGCAA<br>CGGACTCGTT |
| 14451 | CTTGTTGGGC<br>GAACAACCCG  | ATCCGCAAGC<br>TAGGCGTTCC | GGCAACCCTT<br>CCGTTGGGAA  | CCAGGAGGGC<br>GGTCCTCCCG | TTTAGGATCA<br>AAATCCTAGT |
| 14501 | CCTACGATGA<br>GGATGCTACT  | TCTGGAGGGT<br>AGACCTCCCA | GGTAACATTTC<br>CCATTGTAAG | CCGCACTGTT<br>GGCGTGACAA | GGATGTGGAC<br>CCTACACCTG |
| 14551 | GCCTACCAGG<br>CGGATGGTCC  | CGAGCTTGAA<br>GCTCGAACTT | AGATGACACC<br>TCTACTGTGG  | GAACAGGGCG<br>CTTGTCCCGC | GGGGTGGCGC<br>CCCCACCGCG |
| 14601 | AGGCGGCAGC<br>TCCGCCGTCC  | AACAGCAGTG<br>TTGTCGTCAC | GCAGCGGCGC<br>CGTCGCCGCG  | GGAAGAGAAC<br>CCTTCTCTTG | TCCAACGCGG<br>AGGTTGCGCC |
| 14651 | CAGCCGCGGC<br>GTCGGCGCCG  | AATGCAGCCG<br>TTACGTCGGC | GTGGAGGACA<br>CACCTCCTGT  | TGAACGATCA<br>ACTTGCTAGT | TGCCATTTCG<br>ACGGTAAGCG |
| 14701 | GGCGACACCT<br>CCGCTGTGGA  | TTGCCACACG<br>AACGGTGTGC | GGCTGAGGAG<br>CCGACTCCTC  | AAGCGCGCTG<br>TTCGCGCGAC | AGGCCGAAGC<br>TCCGGCTTCG |
| 14751 | AGCGGCCGAA<br>TCGCCGGCTT  | GCTGCCGCCC<br>CGACGGCGGG | CCGCTGCGCA<br>GGCGACGCGT  | ACCCGAGGTC<br>TGGGCTCCAG | GAGAAGCCTC<br>CTCTTCGGAG |
| 14801 | AGAAGAAACC<br>TCTTCTTTGG  | GGTGATCAAA<br>CCACTAGTTT | CCCCTGACAG<br>GGGGACTGTC  | AGGACAGCAA<br>TCCTGTCGTT | GAAACGCAGT<br>CTTTGCGTCA |
| 14851 | TACAACCTAA<br>ATGTTGGATT  | TAAGCAATGA<br>ATTCGTTACT | CAGCACCTTC<br>GTCGTGGAAG  | ACCCAGTACC<br>TGGGTCATGG | GCAGCTGGTA<br>CGTCGACCAT |
| 14901 | CCTTGCCATAC<br>GGAACGTATG | AACTACGGCG<br>TTGATGCCGC | ACCCTCAGAC<br>TGGGAGTCTG  | CGGAATCCGC<br>GCCTTAGGCG | TCATGGACCC<br>AGTACCTGGG |
| 14951 | TGCTTTGCAC<br>ACGAAACGTG  | TCCTGACGTA<br>AGGACTGCAT | ACCTGCGGCT<br>TGGACGCCGA  | CGGAGCAGGT<br>GCCTCGTCCA | CTACTGGTCG<br>GATGACCAGC |
| 15001 | TTGCCAGACA<br>AACGGTCTGT  | TGATGCAAGA<br>ACTACGTTCT | CCCCGTGACC<br>GGGGCACTGG  | TTCCGCTCCA<br>AAGGCGAGGT | CGCGCCAGAT<br>GCGCGGTCTA |
| 15051 | CAGCAACTTT<br>GTCGTTGAAA  | CCGGTGGTGG<br>GGCCACCACC | GCGCCGAGCT<br>CGCGGCTCGA  | GTTGCCCGTG<br>CAACGGGCAC | CACTCCAAGA<br>GTGAGGTTCT |
| 15101 | GCTTCTACAA<br>CGAAGATGTT  | CGACCAGGCC<br>GCTGGTCCGG | GTCTACTCCC<br>CAGATGAGGG  | AACTCATCCG<br>TTGAGTAGGC | CCAGTTTACC<br>GGTCAAATGG |
| 15151 | TCTCTGACCC<br>AGAGACTGGG  | ACGTGTTCAA<br>TGCACAAGTT | TCGCTTTCCC<br>AGCGAAAGGG  | GAGAACCAGA<br>CTCTTGGTCT | TTTTGGCGCG<br>AAAACCGCGC |
| 15201 | CCCGCCAGCC<br>GGGCGGTCGG  | CCCACCATCA<br>GGGTGGTAGT | CCACCGTCAG<br>GGTGGCAGTC  | TGAAAACGTT<br>ACTTTTGCAA | CCTGCTCTCA<br>GGACGAGAGT |

FIG.9A-18

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|       |             |            |            |             |            |
|-------|-------------|------------|------------|-------------|------------|
| 15251 | CAGATCACGG  | GACGCTACCG | CTGCGCAACA | GCATCGGAGG  | AGTCCAGCGA |
|       | GTCTAGTGCC  | CTGCGATGGC | GACGCGTTGT | CGTAGCCTCC  | TCAGGTCGCT |
| 15301 | GTGACCATTA  | CTGACGCCAG | ACGCCGCACC | TGCCCCCTACG | TTTACAAGGC |
|       | CACTGGTAAT  | GACTGCGGTC | TGCGGCGTGG | ACGGGGATGC  | AAATGTTCCG |
| 15351 | CCTGGGCATA  | GTCTCGCCGC | GCGTCCTATC | GAGCCGCACT  | TTTTGAGCAA |
|       | GGACCCGTAT  | CAGAGCGGCG | CGCAGGATAG | CTCGGCGTGA  | AAAACTCGTT |
| 15401 | GCATGTCCAT  | CCTTATATCG | CCCAGCAATA | ACACAGGCTG  | GGGCCTGCGC |
|       | CGTACAGGTA  | GGAATATAGC | GGGTCGTTAT | TGTGTCCGAC  | CCCGGACGCG |
| 15451 | TTCCCAAGCA  | AGATGTTTGG | CGGGGCCAAG | AAGCGCTCCG  | ACCAACACCC |
|       | AAGGGTTCGT  | TCTACAAACC | GCCCCGGTTC | TTCGCGAGGC  | TGGTTGTGGG |
| 15501 | AGTGCGCGTG  | CGCGGGCACT | ACGCGCGGCC | CTGGGGCGCG  | CACAAACGCG |
|       | TCACGCGCAC  | GCGCCCGTGA | TGGCGCGCGG | GACCCCGCGC  | GTGTTTGCGC |
| 15551 | GCCGCACTGG  | GCGCACCACC | GTCGATGACG | CCATCGACGC  | GGTGGTGGAG |
|       | CGGCGTGACC  | CGCGTGGTGG | CAGCTACTGC | GGTAGCTGCG  | CCACCACCTC |
| 15601 | GAGGCGCGCA  | ACTACACGCC | CACGCCGCCA | CCAGTGTCCA  | CAGTGGACGC |
|       | CTCCGCGCGT  | TGATGTGCGG | GTGCGGCGGT | GGTCACAGGT  | GTCACCTGCG |
| 15651 | GGCCATTTCAG | ACCGTGGTGC | GCGGAGCCCG | GCGCTATGCT  | AAAATGAAGA |
|       | CCGGTAAGTC  | TGGCACCACG | CGCCTCGGGC | CGCGATACGA  | TTTTACTTCT |
| 15701 | GACGGCGGAG  | GCGCGTAGCA | CGTCGCCACC | GCCGCCGACC  | CGGCACTGCC |
|       | CTGCCGCCCTC | CGCGCATCGT | GCAGCGGTGG | CGGCGGCTGG  | GCCGTGACGG |
| 15751 | GCCCAACGCG  | CGGCGGCGGC | CCTGCTTAAC | CGCGCACGTC  | GCACCGGCCG |
|       | CGGGTTGCGC  | GCCGCCGCCG | GGACGAATTG | GCGCGTGCA   | CGTGGCCGGC |
| 15801 | ACGGGCGGCC  | ATGCGGGCCG | CTCGAAGGCT | GGCCGCGGGT  | ATTGTCACTG |
|       | TGCCCCGCCG  | TACGCCCGGC | GAGCTTCCGA | CCGGCGCCCA  | TAACAGTGAC |
| 15851 | TGCCCCCAG   | GTCCAGGCGA | CGAGCGGCCG | CCGCAGCAGC  | CGCGGCCATT |
|       | ACGGGGGGTC  | CAGGTCCGCT | GCTCGCCGGC | GGCGTCGTG   | GCGCCGGTAA |
| 15901 | AGTGCTATGA  | CTCAGGGTCG | CAGGGGCAAC | GTGTATTGGG  | TGCGCGACTC |
|       | TCACGATACT  | GAGTCCCAGC | GTCCCCGTTG | CACATAACCC  | ACGCGCTGAG |
| 15951 | GGTTAGCGGC  | CTGCGCGTGC | CCGTGCGCAC | CCGCCCCCG   | CGCAACTAGA |
|       | CCAATCGCCG  | GACGCGCACG | GGCACGCGTG | GGCGGGGGGC  | GCGTTGATCT |
| 16001 | TTGCAAGAAA  | AAACTACTTA | GACTCGTACT | GTTGTATGTA  | TCCAGCGGCG |
|       | AACGTTCTTT  | TTTGATGAAT | CTGAGCATGA | CAACATACAT  | AGGTCGCCGC |
| 16051 | GCGGCGCGCA  | ACGAAGCTAT | GTCCAAGCGC | AAAATCAAAG  | AAGAGATGCT |
|       | CGCCGCGCGT  | TGCTTCGATA | CAGGTTGCGC | TTTTAGTTTC  | TTCTCTACGA |

FIG.9A-19

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|       |            |            |            |            |            |
|-------|------------|------------|------------|------------|------------|
| 16101 | CCAGGTCATC | GCGCCGGAGA | TCTATGGCCC | CCCGAAGAAG | GAAGAGCAGG |
|       | GGTCCAGTAG | CGCGGCCTCT | AGATACCGGG | GGGCTTCTTC | CTTCTCGTCC |
| 16151 | ATTACAAGCC | CCGAAAGCTA | AAGCGGGTCA | AAAAGAAAAA | GAAAGATGAT |
|       | TAATGTTCGG | GGCTTTCGAT | TTCGCCCAGT | TTTTCTTTTT | CTTTCTACTA |
| 16201 | GATGATGAAC | TTGACGACGA | GGTGGAAGT  | CTGCACGCTA | CCGCGCCCAG |
|       | CTACTACTTG | AACTGCTGCT | CCACCTTGAC | GACGTGCGAT | GGCGCGGGTC |
| 16251 | GCGACGGGTA | CAGTGGAAAG | GTCGACGCGT | AAAACGTGTT | TTGCGACCCG |
|       | CGCTGCCCAT | GTCACCTTTC | CAGCTGCGCA | TTTTGCACAA | AACGCTGGGC |
| 16301 | GCACCACCGT | AGTCTTTACG | CCCGGTGAGC | GCTCCACCCG | CACCTACAAG |
|       | CGTGGTGGCA | TCAGAAATGC | GGGCCACTCG | CGAGGTGGGC | GTGGATGTTC |
| 16351 | CGCGTGTATG | ATGAGGTGTA | CGGCGACGAG | GACCTGCTTG | AGCAGGCCAA |
|       | GCGCACATAC | TACTCCACAT | GCCGCTGCTC | CTGGACGAAC | TCGTCCGGTT |
| 16401 | CGAGCGCCTC | GGGGAGTTTG | CCTACGGAAA | GCGGCATAAG | GACATGCTGG |
|       | GCTCGCGGAG | CCCCTCAAAC | GGATGCCTTT | CGCCGTATTC | CTGTACGACC |
| 16451 | CGTTGCCGCT | GGACGAGGGC | AACCCAACAC | CTAGCCTAAA | GCCCCTAACA |
|       | GCAACGGCGA | CCTGCTCCCG | TTGGGTTGTG | GATCGGATTT | CGGGCATTGT |
| 16501 | CTGCAGCAGG | TGCTGCCCGC | GCTTGACCCG | TCCGAAGAAA | AGCGCGGCCT |
|       | GACGTGCTCC | ACGACGGGCG | CGAACGTGGC | AGGCTTCTTT | TCGCGCCGGA |
| 16551 | AAAGCGCGAG | TCTGGTGAAT | TGGCACCCAC | CGTGCAGCTG | ATGGTACCCA |
|       | TTTCGCGCTC | AGACCACTGA | ACCGTGGGTG | GCACGTGAC  | TACCATGGGT |
| 16601 | AGCGCCAGCG | ACTGGAAGAT | GTCTTGGAAT | AAATGACCGT | GGAACCTGGG |
|       | TCGCGGTCGC | TGACCTTCTA | CAGAACCTTT | TTTACTGGCA | CCTTGACCC  |
| 16651 | CTGGAGCCCG | AGGTCCGCGT | GCGGCCAATC | AAGCAGGTGG | CGCCGGGACT |
|       | GACCTCGGGC | TCCAGGCGCA | CGCCGGTTAG | TTCGTCCACC | GCGGCCCTGA |
| 16701 | GGGCGTGACG | ACCGTGGACG | TTCAGATACC | CACTACCACT | AGCACCAGTA |
|       | CCCGCACGTC | TGGCACCTGC | AAGTCTATGG | GTGATGGTCA | TCGTGGTCAT |
| 16751 | TTGCCACCGC | CACAGAGGGC | ATGGAGACAC | AAACGTCCCC | GGTTGCCTCA |
|       | AACGGTGGCG | GTGTCTCCCG | TACCTCTGTG | TTTGCAGGGG | CCAACGGAGT |
| 16801 | GCGGTGGCGG | ATGCCGCGGT | GCAGGCGGTC | GCTGCGGCCG | CGTCCAAGAC |
|       | CGCCACCGCC | TACGGCGCCA | CGTCCGCCAG | CGACGCCGGC | GCAGGTTCTG |
| 16851 | CTCTACGGAG | GTGCAAACGG | ACCCGTGGAT | GTTTCGCGTT | TCAGCCCCCC |
|       | GAGATGCCTC | CACGTTTGCC | TGGGCACCTA | CAAAGCGCAA | AGTCGGGGGG |
| 16901 | GGCGCCCGCG | CCGTTCGAGG | AAGTACGGCG | CCGCCAGCGC | GCTACTGCCC |
|       | CCGCGGGGCG | GGCAAGCTCC | TTCATGCCGC | GGCGGTCGCG | CGATGACGGG |

FIG.9A-20

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16951 GAATATGCCC TACATCCTTC CATTGCGCCT ACCCCCGGCT ATCGTGGCTA  
 CTTATACGGG ATGTAGGAAG GTAACGCGGA TGGGGGCCGA TAGCACCGAT  
 17001 CACCTACCGC CCCAGAAGAC GAGCAACTAC CCGACGCCGA ACCACCACTG  
 GTGGATGGCG GGGTCTTCTG CTCGTTGATG GGCTGCGGCT TGGTGGTGAC  
 17051 GAACCCGCCG CCGCCGTGCG CGTCGCCAGC CCGTGCTGGC CCCGATTTCC  
 CTTGGGCGGC GCGGCAGCG GCAGCGGTCTG GGCACGACCG GGGCTAAAGG  
 17101 GTGCGCAGGG TGGCTCGCGA AGGAGGCAGG ACCCTGGTGC TGCCAACAGC  
 CACGCGTCCC ACCGAGCGCT TCCTCCGTCC TGGGACCACG ACGGTTGTCTG  
 17151 GCGCTACCAC CCCAGCATCG TTTAAAGCC GGTCTTTGTG GTTCTTGCA  
 CGCGATGGTG GGGTCGTAGC AAATTTTCGG CCAGAAACAC CAAGAACGTC  
 17201 ATATGGCCCT CACCTGCCGC CTCCGTTTCC CGGTGCCGGG ATTCCGAGGA  
 TATACGGGA GTGGACGGCG GAGGCAAAGG GCCACGGCCC TAAGGCTCCT  
 17251 AGAATGCACC GTAGGAGGGG CATGGCCGGC CACGGCCTGA CGGGCGGCAT  
 TCTTACGTGG CATCCTCCCC GTACCGGCCG GTGCCGGACT GCCCGCCGTA  
 17301 GCGTCGTGCG CACCACCGGC GCGGCGCGC GTCGCACCGT CGCATGCGCG  
 CGCAGCACGC GTGGTGGCCG CCGCCGCGCG CAGCGTGGCA GCGTACGCGC  
 17351 GCGGTATCCT GCCCCTCCTT ATTCCACTGA TCGCCGCGGC GATTGGCGCC  
 CGCCATAGGA CGGGGAGGAA TAAGGTGACT AGCGGCGCCG CTAACCGCGG  
 17401 GTGCCCCGAA TTGCATCCGT GGCCTTGCA GCGCAGAGAC ACTGATTAAA  
 CACGGGCCCT AACGTAGGCA CCGGAACGTC CGCGTCTCTG TGACTAATTT  
 17451 AACAAAGTTGC ATGTGGA AAAA ATCAAAATAA AAAGTCTGGA CTCTCACGCT  
 TTGTTCAACG TACACCTTTT TAGTTTTATT TTTCAGACCT GAGAGTGCAG  
 17501 CGCTTGGTCC TGTAATATT TTGTAGAATG GAAGACATCA ACTTTGCGTC  
 GCGAACCAGG ACATTGATAA AACATCTTAC CTTCTGTAGT TGAAACGCAG  
 17551 TCTGGCCCCG CGACACGGCT CGCGCCCGTT CATGGGAAAC TGGCAAGATA  
 AGACCGGGGC GCTGTGCCGA GCGCGGGCAA GTACCCTTTG ACCGTTCTAT  
 17601 TCGGCACCAG CAATATGAGC GGTGGCGCCT TCAGCTGGGG CTCGCTGTGG  
 AGCCGTGGTC GTTATACTCG CCACCGCGGA AGTCGACCCC GAGCGACACC  
 17651 AGCGGCATTA AAAATTTTCGG TTCCACCGTT AAGAACTATG GCAGCAAGGC  
 TCGCCGTAAT TTTTAAAGCC AAGGTGGCAA TTCTTGATAC CGTCGTTCCG  
 17701 CTGGAACAGC AGCACAGGCC AGATGCTGAG GGATAAGTTG AAAGAGCAAA  
 GACCTTGTCG TCGTGTCCGG TCTACGACTC CCTATTCAAC TTTCTCGTTT  
 17751 ATTTCCAACA AAAGGTGGTA GATGGCCTGG CCTCTGGCAT TAGCGGGGTG  
 TAAAGTTGT TTTCCACCAT CTACCGGACC GGAGACCGTA ATCGCCCCAC

FIG.9A-21

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17801 GTGGACCTGG CCAACCAGGC AGTGCAAAAT AAGATTAACA GTAAGCTTGA  
 CACCTGGACC GGTGGGTCCG TCACGTTTAA TTCTAATTGT CATTGCAACT  
 17851 TCCCCGCCCT CCCGTAGAGG AGCCTCCACC GGCCGTGGAG ACAGTGTCTC  
 AGGGGCGGGA GGGCATCTCC TCGGAGGTGG CCGGCACCTC TGTCACAGAG  
 17901 CAGAGGGGCG TGGCGAAAAG CGTCCGCGCC CCGACAGGGA AGAAACTCTG  
 GTCTCCCCGC ACCGCTTTTC GCAGGCGCGG GGCTGTCCCT TCTTTGAGAC  
 17951 GTGACGCAAA TAGACGAGCC TCCCTCGTAC GAGGAGGCAC TAAAGCAAGG  
 CACTGCGTTT ATCTGCTCGG AGGGAGCATG CTCCTCCGTG ATTTGTTCC  
 18001 CCTGCCCACC ACCCGTCCCA TCGCGCCCAT GGCTACCGGA GTGCTGGGCC  
 GGACGGGTGG TGGCAGGGT AGCGCGGGTA CCGATGGCCT CACGACCCGG  
 18051 AGCACACACC CGTAACGCTG GACCTGCCTC CCCCCGCCGA CACCCAGCAG  
 TCGTGTGTGG GCATTGCGAC CTGGACGGAG GGGGGCGGCT GTGGGTCTGC  
 18101 AAACCTGTGC TGCCAGGCCG GACCGCCGTT GTTGTAACCC GTCCTAGCCG  
 TTTGGACACG ACGGTCCGGG CTGGCGGCAA CAACATTGGG CAGGATCGGC  
 18151 CGCGTCCCTG CGCCGCGCCG CCAGCGGTCC GCGATCGTTG CGGCCCCGTAG  
 GCGCAGGGAC GCGGCGCGGC GGTCGCCAGG CGCTAGCAAC GCCGGGCATC  
 18201 CCAGTGGCAA CTGGCAAAGC AACTGAACA GCATCGTGGG TCTGGGGGTG  
 GGTCAACGTT GACCGTTTCG TGTGACTTGT CGTAGCACCC AGACCCCCAC  
 18251 CAATCCCTGA AGCGCCGACG ATGCTTCTGA TAGCTAACGT GTCGTATGTG  
 GTTAGGGACT TCGCGGCTGC TACGAAGACT ATCGATTGCA CAGCATACAC  
 18301 TGTCATGTAT GCGTCCATGT CGCCGCCAGA GGAGCTGCTG AGCCGCCGCG  
 ACAGTACATA CGCAGGTACA GCGGCGGTCT CCTCGACGAC TCGGCGGGCG  
 18351 CGCCCGCTTT CCAAGATGGC TACCCCTTCG ATGATGCCGC AGTGGTCTTA  
 GCGGGCGAAA GGTTCCTACCG ATGGGGAAGC TACTACGGCG TCACCAGAAT  
 18401 CATGCACATC TCGGGCCAGG ACGCCTCGGA GTACCTGAGC CCCGGGCTGG  
 GTACGTGTAG AGCCCGGTCC TCGGAGCCT CATGGACTCG GGGCCCGACC  
 18451 TGCAGTTTGC CCGCGCCACC GAGACGTACT TCAGCCTGAA TAACAAGTTT  
 ACGTCAAACG GCGCGGTGG CTCTGCATGA AGTCGGAATT ATTGTTCAAA  
 18501 AGAAACCCCA CCGTGGCGCC TACGCACGAC GTGACCACAG ACCGGTCCCA  
 TCTTTGGGGT GCCACCGCGG ATGCGTGCTG CACTGGTGTC TGGCCAGGGT  
 18551 GCGTTTGACG CTGCGGTTCA TCCCTGTGGA CCGTGAGGAT ACTGCGTACT  
 CGCAAACCTG GACGCCAAGT AGGGACACCT GGCACCTCTA TGACGCATGA  
 18601 CGTACAAGGC GCGGTTACCC CTAGCTGTGG GTGATAACCG TGTGCTGGAC  
 GCATGTTCCG CGCCAAGTGG GATCGACACC CACTATTGGC ACACGACCTG

FIG.9A-22

31/70

18651 ATGGCTTCCA CGTACTTTGA CATCCGCGGC GTGCTGGACA GGGGCCCTAC  
TACCGAAGGT GCATGAACT GTAGCGCCG CACGACCTGT CCCCGGGATG

18701 TTTTAAGCCC TACTCTGGCA CTGCCTACAA CGCCCTGGCT CCAAGGGTG  
AAAATTCGGG ATGAGACCGT GACGGATGTT GCGGGACCGA GGGTTCCAC

18751 CCCCAAATCC TTGCGAATGG GATGAAGCTG CTA CTGCTCT TGAAATAAAC  
GGGGTTTAGG AACGCTTACC CTACTTCGAC GATGACGAGA ACTTTATTG

18801 CTAGAAGAAG AGGACGATGA CAACGAAGAC GAAGTAGACG AGCAAGCTGA  
GATCTTCTTC TCCTGCTACT GTTGCTTCTG CTTTCATCTGC TCGTTGCACT

18851 GCAGCAAAAA ACTCACGTAT TTGGGCAGGC GCCTTATTCT GGTATAAATA  
CGTCGTTTTT TGAGTGCATA AACCCGTCCG CGGAATAAGA CCATATTTAT

18901 TTACAAAGGA GGGTATTCAA ATAGGTGTCG AAGGTCAAAC ACCTAAATAT  
AATGTTTCCT CCCATAAGTT TATCCACAGC TTCCAGTTTG TGGATTTATA

18951 GCCGATAAAA CATTTCAACC TGAACCTCAA ATAGGAGAAT CTCAGTGGA  
CGGCTATTTT GTAAAGTTGG ACTTGAGTT TATCCTCTTA GAGTCACCAT

19001 CGAAACAGAA ATTAATCATG CAGCTGGGAG AGTCCTAAAA AAGACTACCC  
GCTTTGTCTT TAATTAGTAC GTCGACCCTC TCAGGATTTT TTCTGATGGG

19051 CAATGAAACC ATGTTACGGT TCATATGCAA AACCCACAAA TGAAAATGGA  
GTTACTTTGG TACAATGCCA AGTATACGTT TTGGGTGTTT ACTTTTACCT

19101 GGGCAAGGCA TTCTTGTAAG GCAACAAAAT GGAAAGCTAG AAAGTCAAGT  
CCCGTTCCGT AAGAACATTT CGTTGTTTTA CCTTTCGATC TTTCAGTTCA

19151 GGAAATGCAA TTTTCTCAA CTA CTGAGGC AGCCGCAGGC AATGGTGATA  
CCTTTACGTT AAAAAGAGTT GATGACTCCG TCGGCGTCCG TTACCACTAT

19201 ACTTGACTCC TAAAGTGGA TTGTACAGTG AAGATGTAGA TATAGAAACC  
TGAAGTGAAG ATTTACCAT AACATGTCAC TTCTACATCT ATATCTTTGG

19251 CCAGACACTC ATATTTCTTA CATGCCCACT ATTAAGGAAG GTA ACTCACG  
GGTCTGTGAG TATAAAGAAT GTACGGGTGA TAATTCCTTC CATTGAGTGC

19301 AGAACTAATG GGCCAACAAT CTATGCCCAA CAGGCCTAAT TACATTGCTT  
TCTTGATTAC CCGGTTGTTA GATACGGGT GTCCGGATTA ATGTAACGAA

19351 TTAGGGACAA TTTTATTGGT CTAATGTATT ACAACAGCAC GGGTAATATG  
AATCCCTGTT AAAATAACCA GATTACATAA TGTTGTCGTG CCCATTATAC

19401 GGTGTTCTGG CGGGCCAAGC ATCGCAGTTG AATGCTGTTG TAGATTTGCA  
CCACAAGACC GCCCGGTTTC TAGCGTCAAC TTACGACAAC ATCTAAACGT

19451 AGACAGAAAC ACAGAGCTTT CATAACAGCT TTTGCTTGAT TCCATTGGTG  
TCTGTCTTTG TGTCTCGAAA GTATGGTCGA AAACGAAC TA AGGTAACCA

FIG.9A-23

32/70

19501 ATAGAACCAG GTACTTTTCT ATGTGGAATC AGGCTGTTGA CAGCTATGAT  
TATCTTGGTC CATGAAAAGA TACACCTTAG TCCGACAACT GTCGATACTA

19551 CCAGATGTTA GAATTATTGA AAATCATGGA ACTGAAGATG AACTTCCAAA  
GGTCTACAAT CTTAATAACT TTTAGTACCT TGAATTCTAC TTGAAGGTTT

19601 TTAAGTCTTT CCACTGGGAG GTGTGATTAA TACAGAGACT CTTACCAAGG  
AATGACGAAA GGTGACCCTC CACACTAATT ATGTCTCTGA GAATGGTTCC

19651 TAAAACCTAA AACAGGTCAG GAAAATGGAT GGGAAAAAGA TGCTACAGAA  
ATTTTGGATT TTGTCCAGTC CTTTACCTA CCCTTTTTCT ACAGTGTCTT

19701 TTTTCAGATA AAAATGAAAT AAGAGTTGGA AATAATTTTG CCATGGAAAT  
AAAAGTCTAT TTTTACTTTA TTCTCAACCT TTATTAAAAC GGTACCTTTA

19751 CAATCTAAAT GCCAACCTGT GGAGAAATTT CCTGTACTCC AACATAGCGC  
GTTAGATTTA CGGTTGGACA CCTCTTTAAA GGACATGAGG TTGTATCGCG

19801 TGTATTTGCC CGACAAGCTA AAGTACAGTC CTTCCAACGT AAAAATTTCT  
ACATAAACGG GCTGTTTCAT TTCATGTCAG GAAGGTTGCA TTTTAAAGA

19851 GATAACCCAA ACACCTACGA CTACATGAAC AAGCGAGTGG TGGCTCCCGG  
CTATTGGGTT TGTGGATGCT GATGTACTTG TTCGCTCACC ACCGAGGGCC

19901 GCTAGTGGAC TGCTACATTA ACCTTGGAGC ACGCTGGTCC CTTGACTATA  
CGATCACCTG ACGATGTAAT TGGAACCTCG TGCGACCAGG GAACTGATAT

19951 TGGACAACGT CAACCCATTT AACCACCACC GCAATGCTGG CCTGCGCTAC  
ACCTGTTGCA GTTGGGTAAA TTGGTGGTGG CGTTACGACC GGACGCGATG

20001 CGCTCAATGT TGCTGGGCAA TGGTCGCTAT GTGCCCTTCC ACATCCAGGT  
GCGAGTTACA ACGACCCGTT ACCAGCGATA CACGGGAAGG TGTAGGTCCA

20051 GCCTCAGAAG TTCTTTGCCA TAAAAACCT CTTTCTCCTG CCGGGCTCAT  
CGGAGTCTTC AAGAAACGGT AATTTTTGGA GGAAGAGGAC GGCCCAGTA

20101 ACACCTACGA GTGGAACCTC AGGAAGGATG TTAACATGGT TCTGCAGAGC  
TGTGGATGCT CACCTTGAAG TCCTTCCTAC AATTGTACCA AGACGTCTCG

20151 TCCCTAGGAA ATGACCTAAG GGTGACGGA GCCAGCATT AATTTGATAG  
AGGGATCCTT TACTGGATTC CCAACTGCCT CGGTCGTAAT TCAAATATC

20201 CATTTGCCTT TACGCCACCT TCTTCCCAT GGCCACAAC ACCGCCTCCA  
GTAAACGGAA ATGCGGTGGA AGAAGGGGTA CCGGGTGTTG TGGCGGAGGT

20251 CGCTTGAGGC CATGCTTAGA AACGACACCA ACGACCAGTC CTTTAACGAC  
GCGAACTCCG GTACGAATCT TTGCTGTGGT TGCTGGTCAG GAAATTGCTG

20301 TATCTCTCCG CCGCCAACAT GCTCTACCCT ATACCCGCCA ACGCTACCAA  
ATAGAGAGGC GGCAGTTGTA CGAGATGGGA TATGGGCGGT TGCGATGGTT

FIG.9A-24



33/70

20351 CGTGCCCAT TCCATCCCCT CCCGCAACTG GGCGGCTTTC CGCGGCTGGG  
 GCACGGGTAT AGGTAGGGGA GGGCGTTGAC CCGCCGAAAG GCGCCGACCC  
 20401 CCTTCACGCG CCTTAAGACT AAGGAAACCC CATCACTGGG CTCGGGCTAC  
 GGAAGTGCGC GGAATTCTGA TTCCTTTGGG GTAGTGACCC GAGCCCGATG  
 20451 GACCCTTATT ACACCTACTC TGGCTCTATA CCCTACCTAG ATGGAACCTT  
 CTGGGAATAA TGTGGATGAG ACCGAGATAT GGGATGGATC TACCTTGGAA  
 20501 TTACCTCAAC CACACCTTTA AGAAGGTGGC CATTACCTTT GACTCTTCTG  
 AATGGAGTTG GTGTGGAAAT TCTCCACCG GTAATGGAAA CTGAGAAGAC  
 20551 TCAGCTGGCC TGGCAATGAC CGCCTGCTTA CCCCCAACGA GTTTGAAATT  
 AGTCGACCGG ACCGTTACTG GCGGACGAAT GGGGGTTGCT CAAACTTTAA  
 20601 AAGCGCTCAG TTGACGGGGA GGGTTACAAC GTTGCCCAGT GTAACATGAC  
 TTCGCGAGTC AACTGCCCCT CCCAATGTTG CAACGGGTCA CATTGTACTG  
 20651 CAAAGACTGG TTCCTGGTAC AAATGCTAGC TAACTATAAC ATTGGCTACC  
 GTTTCTGACC AAGGACCATG TTTACGATCG ATTGATATTG TAACCGATGG  
 20701 AGGGCTTCTA TATCCCAGAG AGCTACAAGG ACCGCATGTA CTCCTTCTTT  
 TCCCGAAGAT ATAGGGTCTC TCGATGTTCC TGGCGTACAT GAGGAAGAAA  
 20751 AGAAACTTCC AGCCCATGAG CCGTCAGGTG GTGGATGATA CTAAATACAA  
 TCTTTGAAGG TCGGGTACTC GGCAGTCCAC CACCTACTAT GATTTATGTT  
 20801 GGACTIONCAA CAGGTGGGCA TCCTACACCA ACACAACAAC TCTGGATTTG  
 CCTGATGGTT GTCCACCCGT AGGATGTGGT TGTGTTGTTG AGACCTAAAC  
 20851 TTGGCTACCT TGCCCCCACC ATGCGCGAAG GACAGGCCTA CCCTGCTAAC  
 AACCAGTGA ACGGGGGTGG TACGCGCTTC CTGTCCGGAT GGGACGATTG  
 20901 TTCCCCTATC CGCTTATAGG CAAGACCGCA GTTGACAGCA TTACCCAGAA  
 AAGGGGATAG GCGAATATCC GTTCTGGCGT CAACTGTCGT AATGGGTCTT  
 20951 AAAGTTTCTT TGCATCGCA CCCTTTGGCG CATCCCATTC TCCAGTAACT  
 TTTCAAAGAA ACGCTAGCGT GGGAAACCGC GTAGGGTAAG AGGTCATTGA  
 21001 TTATGTCCAT GGGCGCACTC ACAGACCTGG GCCAAAACCT TCTCTACGCC  
 AATACAGGTA CCCGCGTGAG TGTCTGGACC CGGTTTTGGA AGAGATGCGG  
 21051 AACTCCGCCC ACGCGCTAGA CATGACTTTT GAGGTGGATC CCATGGACGA  
 TTGAGGCGGG TGC GCGATCT GTACTGAAAA CTCCACCTAG GGTACCTGCT  
 21101 GCCCACCCTT CTTTATGTTT TGTTTGAAGT CTTTGACGTG GTCCGTGTGC  
 CGGGTGGGAA GAAATACAAA ACAAACTTCA GAAACTGCAC CAGGCACACG  
 21151 ACCAGCCGCA CCGCGGCGTC ATCGAAACCG TGTACCTGCG CACGCCCTTC  
 TGGTCGGCGT GGCGCCGCG TAGCTTTGGC ACATGGACGC GTGCGGGGAAG

FIG.9A-25

34/70

21201 TCGGCCGGCA ACGCCACAAC ATAAAGAAGC AAGCAACATC AACAACAGCT  
 AGCCGGCCGT TGCAGTGTG TATTCTTCG TTCGTTGTAG TTGTTGTCTGA  
 21251 GCCGCCATGG GCTCCAGTGA GCAGGAACTG AAAGCCATTG TCAAAGATCT  
 CGGCGGTACC CGAGGTCACT CGTCCTTGAC TTTCGGTAAC AGTTTCTAGA  
 21301 TGGTTGTGGG CCATATTTTT TGGGCACCTA TGACAAGCGC TTTCAGGCT  
 ACCAACACCC GGTATAAAAA ACCCGTGGAT ACTGTTTCGCG AAAGGTCCGA  
 21351 TTGTTTCTCC ACACAAGCTC GCCTGCGCCA TAGTCAATAC GGCCGGTCGC  
 AACAAAGAGG TGTGTTTCGAG CGGACGCGGT ATCAGTTATG CCGGCCAGCG  
 21401 GAGACTGGGG GCGTACACTG GATGGCCTTT GCCTGGAACC CGCACTCAAA  
 CTCTGACCCC CGCATGTGAC CTACCGGAAA CGGACCTTGG GCGTGAGTTT  
 21451 AACATGCTAC CTCTTTGAGC CCTTTGGCTT TTCTGACCAG CGACTCAAGC  
 TTGTACGATG GAGAACTCG GGAAACCGAA AAGACTGGTC GCTGAGTTCTG  
 21501 AGGTTTACCA GTTTGAGTAC GAGTCACTCC TGCGCCGTAG CGCCATTGCT  
 TCCAAATGGT CAAACTCATG CTCAGTGAGG ACGCGGCATC GCGGTAACGA  
 21551 TCTTCCCCCG ACCGCTGTAT AACGCTGGAA AAGTCCACCC AAAGCGTACA  
 AGAAGGGGGC TGGCGACATA TTGCGACCTT TTCAGGTGGG TTTCGCATGT  
 21601 GGGGCCCAAC TCGGCCGCCT GTGGACTATT CTGCTGCATG TTTCTCCACG  
 CCCCAGGTTG AGCCGGCGGA CACCTGATAA GACGACGTAC AAAGAGGTGC  
 21651 CCTTTGCCAA CTGGCCCCAA ACTCCCATGG ATCACAACCC CACCATGAAC  
 GGAAACGGTT GACCGGGGTT TGAGGGTACC TAGTGTTGGG GTGGTACTTG  
 21701 CTTATTACCG GGGTACCCAA CTCCATGCTC AACAGTCCCC AGGTACAGCC  
 GAATAATGGC CCCATGGGTT GAGGTACGAG TTGTCAGGGG TCCATGTCGG  
 21751 CACCCTGCGT CGCAACCAGG AACAGCTCTA CAGCTTCCTG GAGCGCCACT  
 GTGGGACGCA GCGTTGGTCC TTGTCGAGAT GTCGAAGGAC CTCGCGGTGA  
 21801 CGCCCTACTT CCGCAGCCAC AGTGCGCAGA TTAGGAGCGC CACTTCTTTT  
 GCGGGATGAA GCGTCGGTG TCACGCGTCT AATCCTCGCG GTGAAGAAAA  
 21851 TGTCACCTGA AAAACATGTA AAAATAATGT ACTAGAGACA CTTTCAATAA  
 ACAGTGAAC TTTTGTACAT TTTTATTACA TGATCTCTGT GAAAGTTATT  
 21901 AGGCAAATGC TTTTATTTGT AACTCTCGG GTGATTATTT ACCCCCACCC  
 TCCGTTTACG AAAATAAACA TGTGAGAGCC CACTAATAAA TGGGGGTGGG  
 21951 TTGCCGTCTG CGCCGTTTAA AAATCAAAGG GGTTCTGCCG CGCATCGCTA  
 AACGGCAGAC GCGGCAAATT TTTAGTTTCC CCAAGACGGC GCGTAGCGAT  
 22001 TGCGCCACTG GCAGGGACAC GTTGCGATAC TGGTGTTTAG TGCTCCACTT  
 ACGCGGTGAC CGTCCCTGTG CAACGCTATG ACCACAAATC ACGAGGTGAA

FIG.9A-26

35/70

22051 AAACTCAGGC ACAACCATCC GCGGCAGCTC GGTGAAGTTT TCACTCCACA  
 TTTGAGTCCG TGTTGGTAGG CGCCGTCGAG CCACTTCAAA AGTGAGGTGT  
 22101 GGCTGCGCAC CATCACCAAC GCGTTTAGCA GGTGCGGCGC CGATATCTTG  
 CCGACGCGTG GTAGTGGTTG CGCAAATCGT CCAGCCCGCG GCTATAGAAC  
 22151 AAGTCGCAGT TGGGGCCTCC GCCCTGCGCG CGCGAGTTGC GATACACAGG  
 TTCAGCGTCA ACCCCGGAGG CGGGACGCGC GCGCTCAACG CTATGTGTCC  
 22201 GTTGCGAGCAC TGGAACACTA TCAGCGCCGG GTGGTGCACG CTGGCCAGCA  
 CAACGTCGTG ACCTTGTGAT AGTCGCGGCC CACCACGTGC GACCGGTCGT  
 22251 CGCTCTTGTC GGAGATCAGA TCCGCGTCCA GGTCTCCGC GTTGCTCAGG  
 GCGAGAACAG CCTCTAGTCT AGGCGCAGGT CCAGGAGGCG CAACGAGTCC  
 22301 GCGAACGGAG TCAACTTTGG TAGCTGCCTT CCCAAAAGG GCGCGTGCCC  
 CGTTGCCTC AGTTGAAACC ATCGACGGAA GGGTTTTTCC CGCGCACGGG  
 22351 AGGCTTTGAG TTGCACTCGC ACCGTAGTGG CATCAAAGG TGACCGTGCC  
 TCCGAAACTC AACGTGAGCG TGGCATCACC GTAGTTTTCC ACTGGCACGG  
 22401 CGGTCTGGGC GTTAGGATAC AGCGCCTGCA TAAAGCCTT GATCTGCTTA  
 GCCAGACCCG CAATCCTATG TCGCGGACGT ATTTTCGGAA CTAGACGAAT  
 22451 AAAGCCACCT GAGCCTTTGC GCCTTCAGAG AAGAACATGC CGCAAGACTT  
 TTTCGGTGGA CTCGGAACG CGGAAGTCTC TTCTTGACG GCGTTCTGAA  
 22501 GCCGGAAAAC TGATTGGCCG GACAGGCCGC GTCGTGCACG CAGCACCTTG  
 CGGCCTTTTG ACTAACCAGC CTGTCCGGCG CAGCACGTGC GTCGTGGAAC  
 22551 CGTCGGTGTT GGAGATCTGC ACCACATTTT GGCCCCACCG GTTCTTCACG  
 GCAGCCACAA CCTCTAGACG TGGTGTAAG CCGGGGTGGC CAAGAAGTGC  
 22601 ATCTTGGCCT TGCTAGACTG CTCCTTCAGC GCGCGCTGCC CGTTTTCGCT  
 TAGAACCGBA ACGATCTGAC GAGGAAGTCG CGCGCGACGG GCAAAAGCGA  
 22651 CGTCACATCC ATTTCAATCA CGTGCTCCTT ATTTATCATA ATGCTTCCGT  
 GCAGTGTAGG TAAAGTTAGT GCACGAGGAA TAAATAGTAT TACGAAGGCA  
 22701 GTAGACACTT AAGCTCGCCT TCGATCTCAG CGCAGCGGTG CAGCCACAAC  
 CATCTGTGAA TTCGAGCGGA AGCTAGAGTC GCGTCGCCAC GTCGGTGTG  
 22751 GCGCAGCCCG TGGGCTCGTG ATGCTTGTAG GTCACCTCTG CAAACGACTG  
 CGCGTCGGGC ACCCGAGCAC TACGAACATC CAGTGGAGAC GTTTGCTGAC  
 22801 CAGGTACGCC TGCAGGAATC GCCCATCAT CGTCACAAAG GTCTTGTTGC  
 GTCCATGCGG ACGTCCTTAG CGGGGTAGTA GCAGTGTTC CAGAACAACG  
 22851 TGGTGAAGGT CAGCTGCAAC CCGCGGTGCT CCTCGTTCAG CCAGGTCTTG  
 ACCACTTCCA GTCGACGTTG GGCGCCACGA GGAGCAAGTC GTCCAGAAC

FIG.9A-27

36/70

22901 CATACGGCCG CCAGAGCTTC CACTTGGTCA GGCAGTAGTT TGAAGTTCGC  
 GTATGCCGGC GGTCTCGAAG GTGAACCAGT CCGTCATCAA ACTTCAAGCG  
 22951 CTTTAGATCG TTATCCACGT GGTACTTGTC CATCAGCGCG CGCGCAGCCT  
 GAAATCTAGC AATAGGTGCA CCATGAACAG GTAGTCGCGC GCGCGTCGGA  
 23001 CCATGCCCTT CTCCCACGCA GACACGATCG GCACACTCAG CGGGTTCATC  
 GGTACGGGAA GAGGGTGCGT CTGTGCTAGC CGTGTGAGTC GCCCAAGTAG  
 23051 ACCGTAATTT CACTTTCCGC TTCGCTGGGC TCTTCTCTT CCTCTTGCGT  
 TGGCATTAAA GTGAAAGGCG AAGCGACCCG AGAAGGAGAA GGAGAACGCA  
 23101 CCGCATACCA CGCGCCACTG GGTCGTCTTC ATTCAGCCGC CGCACTGTGC  
 GGGGTATGGT GCGCGGTGAC CCAGCAGAAG TAAGTCGGCG GCGTGACACG  
 23151 GCTTACCTCC TTTGCCATGC TTGATTAGCA CCGGTGGGTT GCTGAAACCC  
 CGAATGGAGG AAACGGTACG AACTAATCGT GGCCACCCAA CGACTTTGGG  
 23201 ACCATTTGTA GCGCCACATC TTCTCTTCT TCCTCGCTGT CCACGATTAC  
 TGGTAAACAT CGCGGTGTAG AAGAGAAAGA AGGAGCGACA GGTGCTAATG  
 23251 CTCTGGTGAT GGCGGGCGCT CGGGCTTGGG AGAAGGGCGC TTCTTTTTCT  
 GAGACCACTA CCGCCCGCGA GCCCGAACCC TCTTCCCGCG AAGAAAAAGA  
 23301 TCTTGGGCGC AATGGCCAAA TCCGCCGCCG AGGTCGATGG CCGCGGGCTG  
 AGAACCCGCG TTACCGGTTT AGGCGGCGGC TCCAGCTACC GGCGCCCGAC  
 23351 GGTGTGCGCG GCACCAGCGC GTCTTGATGAT GAGTCTTCCT CGTCCTCGGA  
 CCACACGCGC CGTGGTCGCG CAGAACACTA CTCAGAAGGA GCAGGAGCCT  
 23401 CTCGATACGC CGCCTCATCC GCTTTTTTGG GGGCGCCCGG GGAGGCGGCG  
 GAGCTATGCG GCGGAGTAGG CGAAAAAACC CCCGCGGGCC CCTCCGCCG  
 23451 GCGACGGGGA CGGGGACGAC ACGTCCTCCA TGGTTGGGGG ACGTCGCGCC  
 CGCTGCCCCCT GCCCCTGCTG TGCAGGAGGT ACCAACCCCC TGCAGCGCGG  
 23501 GCACCGCGTC CGCGCTCGGG GGTGGTTTCG CGCTGCTCCT CTTCCCGACT  
 CGTGGCGCAG GCGCGAGCCC CCACCAAAGC GCGACGAGGA GAAGGGCTGA  
 23551 GGCCATTTCC TTCTCCTATA GGCAGAAAAA GATCATGGAG TCAGTCGAGA  
 CCGGTAAAGG AAGAGGATAT CCGTCTTTTT CTAGTACCTC AGTCAGCTCT  
 23601 AGAAGGACAG CCTAACCGCC CCCTCTGAGT TCGCCACCAC CGCCTCCACC  
 TCTTCCTGTC GGATTGGCGG GGGAGACTCA AGCGGTGGTG GCGGAGGTGG  
 23651 GATGCCGCCA ACGCGCCTAC CACCTTCCCC GTCGAGGCAC CCCCCTTGA  
 CTACGGCGGT TGCGCGGATG GTGGAAGGGG CAGCTCCGTG GGGGCGAACT  
 23701 GGAGGAGGAA GTGATTATCG AGCAGGACCC AGGTTTTGTA AGCGAAGACG  
 CCTCCTCCTT CACTAATAGC TCGTCCTGGG TCCAAAACAT TCGCTTCTGC

FIG.9A-28

37/70

23751 ACGAGGACCG CTCAGTACCA ACAGAGGATA AAAAGCAAGA CCAGGACAAC  
 TGCTCCTGGC GAGTCATGGT TGTCTCCTAT TTTTCGTTCT GGTCTGTG

23801 GCAGAGGCAA ACGAGGAACA AGTCGGGCGG GGGGACGAAA GGCATGGCGA  
 CGTCTCCGTT TGCTCCTTGT TCAGCCCGCC CCCCTGCTTT CCGTACCGCT

23851 CTACCTAGAT GTGGGAGACG ACGTGCTGTT GAAGCATCTG CAGCGCCAGT  
 GATGGATCTA CACCCTCTGC TGCACGACAA CTTCGTAGAC GTCGCGGTCA

23901 GCGCCATTAT CTGCGACGCG TTGCAAGAGC GCAGCGATGT GCCCCTCGCC  
 CGCGGTAATA GACGCTGCGC AACGTTCTCG CGTCGCTACA CGGGGAGCGG

23951 ATAGCGGATG TCAGCCTTGC CTACGAACGC CACCTATTCT CACCGCGCGT  
 TATCGCCTAC AGTCGGAACG GATGCTTGCG GTGGATAAGA GTGGCGCGCA

24001 ACCCCCCAAA CGCCAAGAAA ACGGCACATG CGAGCCCAAC CCGCGCCTCA  
 TGGGGGGTTT GCGGTTCTTT TGCCGTGTAC GCTCGGGTTG GGCGCGGAGT

24051 ACTTCTACCC CGTATTTGCC GTGCCAGAGG TGCTTGCCAC CTATCACATC  
 TGAAGATGGG GCATAAACGG CACGGTCTCC ACGAACGGTG GATAGTGTAG

24101 TTTTTCCAAA ACTGCAAGAT ACCCCTATCC TGCCGTGCCA ACCGCAGCCG  
 AAAAAGGTTT TGACGTTCTA TGGGGATAGG ACGGCACGGT TGGCGTCGGC

24151 AGCGGACAAG CAGCTGGCCT TCGGGCAGGG CGCTGTCATA CCTGATATCG  
 TCGCCTGTTC GTCGACCGGA ACGCCGTCCC GCGACAGTAT GGACTATAGC

24201 CCTCGCTCAA CGAAGTGCCA AAAATCTTTG AGGGTCTTGG ACGCGACGAG  
 GGAGCGAGTT GCTTCACGGT TTTTAGAAAC TCCCAGAACC TGCCTGCTC

24251 AAGCGCGCGG CAAACGCTCT GCAACAGGAA AACAGCGAAA ATGAAAGTCA  
 TTCGCGCGCC GTTTGCGAGA CGTTGTCCTT TTGTCGCTTT TACTTTCAGT

24301 CTCTGGAGTG TTGGTGGAAC TCGAGGGTGA CAACGCGCGC CTAGCCGTAC  
 GAGACCTCAC AACCACCTTG AGCTCCCACT GTTGCGCGCG GATCGGCATG

24351 TAAAACGCAG CATCGAGGTC ACCCACTTTG CCTACCCGGC ACTTAACCTA  
 ATTTTGCGTC GTAGCTCCAG TGGGTGAAAC GGATGGGCCG TGAATTGGAT

24401 CCCCCCAAGG TCATGAGCAC AGTCATGAGT GAGCTGATCG TGCGCCGTGC  
 GGGGGGTTCC AGTACTCGTG TCAGTACTCA CTCGACTAGC ACGCGGCACG

24451 GCAGCCCCTG GAGAGGGATG CAAATTTGCA AGAACAAACA GAGGAGGGCC  
 CGTCGGGGAC CTCTCCCTAC GTTTAAACGT TCTTGTTTGT CTCCTCCCGG

24501 TACCCGCAGT TGGCGACGAG CAGCTAGCGC GCTGGCTTCA AACGCGCGAG  
 ATGGGCGTCA ACCGCTGCTC GTCGATCGCG CGACCGAAGT TTGCGCGCTC

24551 CCTGCCGACT TGGAGGAGCG ACGCAAATA ATGATGGCCG CAGTGCTCGT  
 GGACGGCTGA ACCTCCTCGC TGCCTTTGAT TACTACCGGC GTCACGAGCA

FIG.9A-29

38/70

24601 TACCGTGGAG CTTGAGTGCA TGCAGCGGTT CTTTGCTGAC CCGGAGATGC  
 ATGGCACCTC GAACTCACGT ACGTCGCCAA GAAACGACTG GGCCTCTACG  
 24651 AGCGCAAGCT AGAGGAAACA TTGCACTACA CCTTTCGACA GGGCTACGTA  
 TCGCGTTCGA TCTCCTTTGT AACGTGATGT GGAAAGCTGT CCCGATGCAT  
 24701 CGCCAGGCCT GCAAGATCTC CAACGTGGAG CTCTGCAACC TGGTCTCCTA  
 GCGGTCCGGA CGTTCTAGAG GTTGCACCTC GAGACGTTGG ACCAGAGGAT  
 24751 CCTTGGAATT TTGCACGAAA ACCGCCTTGG GCAAAACGTG CTTCAATTCCA  
 GGAACCTTAA AACGTGCTTT TGGCGGAACC CGTTTTGCAC GAAGTAAGGT  
 24801 CGCTCAAGGG CGAGGCGCGC GCGCACTACG TCCGCGACTG CGTTTACTTA  
 GCGAGTTCCC GCTCCGCGCG GCGCTGATGC AGGCGCTGAC GCAAATGAAT  
 24851 TTTCTATGCT ACACCTGGCA GACGGCCATG GGC GTTTGGC AGCAGTGCTT  
 AAAGATACGA TGTGGACCGT CTGCCGGTAC CCGCAAACCG TCGTCACGAA  
 24901 GGAGGAGTGC AACCTCAAGG AGCTGCAGAA ACTGCTAAAG CAAAATTGA  
 CCTCCTCACG TTGGAGTTCC TCGACGTCTT TGACGATTTT GTTTTGAAC  
 24951 AGGACCTATG GACGGCCTTC AACGAGCGCT CCGTGGCCGC GCACCTGGCG  
 TCCTGGATAC CTGCCGGAAG TTGCTCGCGA GGCACCGGCG CGTGGACCGC  
 25001 GACATCATTT TCCCCGAACG CCTGCTTAAA ACCCTGCAAC AGGGTCTGCC  
 CTGTAGTAAA AGGGGCTTGC GGACGAATTT TGGGACGTTG TCCAGACGG  
 25051 AGACTTCACC AGTCAAAGCA TGTTGCAGAA CTTTAGGAAC TTTATCCTAG  
 TCTGAAGTGG TCAGTTTCGT ACAACGTCTT GAAATCCTTG AAATAGGATC  
 25101 AGCGCTCAGG AATCTTGCCC GCCACCTGCT GTGCACTTCC TAGCGACTTT  
 TCGCGAGTCC TTAGAACGGG CCGTGGACGA CACGTGAAGG ATCGCTGAAA  
 25151 GTGCCCATTG AGTACCGCGA ATGCCCTCCG CCGCTTTGGG GCCACTGCTA  
 CACGGGTAAT TCATGGCGCT TACGGGAGGC GGC GAAACCC CCGTGACGAT  
 25201 CCTTCTGCAG CTAGCCAACT ACCTTGCCCTA CCACTCTGAC ATAATGGAAG  
 GGAAGACGTC GATCGGTTGA TGGAACGGAT GGTGAGACTG TATTACCTTC  
 25251 ACGTGAGCGG TGACGGTCTA CTGGAGTGTC ACTGTCGCTG CAACCTATGC  
 TGCACTCGCC ACTGCCAGAT GACCTCACAG TGACAGCGAC GTTGGATACG  
 25301 ACCCCGCACC GCTCCCTGGT TTGCAATTCG CAGCTGCTTA ACGAAAGTCA  
 TGGGGCGTGG CGAGGGACCA AACGTTAAGC GTCGACGAAT TGCTTTCAGT  
 25351 AATTATCGGT ACCTTTGAGC TGCAGGGTCC CTCGCCTGAC GAAAAGTCCG  
 TTAATAGCCA TGGAACCTCG ACGTCCCAGG GAGCGGACTG CTTTTACGGC  
 25401 CGGCTCCGGG GTTGAAACTC ACTCCGGGGC TGTGGACGTC GGCTTACCTT  
 GCCGAGGCC CAACCTTGAG TGAGGCCCG ACACCTGCAG CCGAATGGAA

FIG.9A-30

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25451 CGCAAATTTG TACCTGAGGA CTACCACGCC CACGAGATTA GGTTCCTACGA  
 GCGTTTAAAC ATGGACTCCT GATGGTGCGG GTGCTCTAAT CCAAGATGCT  
 25501 AGACCAATCC CGCCCGCCTA ATGCGGAGCT TACCGCCTGC GTCATTACCC  
 TCTGGTTAGG GCGGGCGGAT TACGCCTCGA ATGGCGGACG CAGTAATGGG  
 25551 AGGGCCACAT TCTTGGCCAA TTGCAAGCCA TCAACAAAGC CCGCCAAGAG  
 TCCCGGTGTA AGAACCGGTT AACGTTGCGT AGTTGTTTCG GCGGGTTCTC  
 25601 TTTCTGCTAC GAAAGGGACG GGGGGTTTAC TTGGACCCCC AGTCCGGCGA  
 AAAGACGATG CTTTCCCTGC CCCCCAAATG AACCTGGGGG TCAGGCCGCT  
 25651 GGAGCTCAAC CCAATCCCCC CGCCGCCGCA GCCCTATCAG CAGCAGCCGC  
 CCTCGAGTTG GGTAGGGGG GCGGCGGCGT CGGGATAGTC GTCGTCGGCG  
 25701 GGGCCCTTGC TTCCCAGGAT GGCACCCAAA AAGAAGCTGC AGCTGCCGCC  
 CCCGGGAACG AAGGGTCCTA CCGTGGGTTT TTCTTCGACG TCGACGGCGG  
 25751 GCCACCCACG GACGAGGAGG AATACTGGGA CAGTCAGGCA GAGGAGGTTT  
 CGGTGGGTGC CTGCTCCTCC TTATGACCCT GTCAGTCCGT CTCCTCCAAA  
 25801 TGGACGAGGA GGAGGAGGAC ATGATGGAAG ACTGGGAGAG CCTAGACGAG  
 ACCTGCTCCT CCTCCTCTG TACTACCTTC TGACCCTCTC GGATCTGCTC  
 25851 GAAGCTTCCG AGGTCGAAGA GGTGTCAGAC GAAACACCGT CACCCTCGGT  
 CTTCGAAGGC TCCAGCTTCT CCACAGTCTG CTTTGTGGCA GTGGGAGCCA  
 25901 CGCATTCCCC TCGCCGGCGC CCCAGAAATC GGCAACCGGT TCCAGCATGG  
 GCGTAAGGGG AGCGGCCGCG GGGTCTTTAG CCGTTGGCCA AGGTCGTACC  
 25951 CTACAACCTC CGCTCCTCAG GCGCCGCCGG CACTGCCCCTG TCGCCGACCC  
 GATGTTGGAG GCGAGGAGTC GCGGGCGGCC GTGACGGGCA AGCGGCTGGG  
 26001 AACCCTAGAT GGGACACCAC TGGAAACCAGG GCCGGTAAGT CCAAGCAGCC  
 TTGGCATCTA CCCTGTGGTG ACCTTGGTCC CGGCCATTCA GGTTGTCGCG  
 26051 GCCGCCGTTA GCCCAAGAGC AACAAACAGC CCAAGGCTAC CGCTCATGGC  
 CGGCGGCAAT CGGGTTCTCG TTGTTGTGCG GGTTCGATG GCGAGTACCG  
 26101 GCGGGCACAA GAACGCCATA GTTGCTTGCT TGCAAGACTG TGGGGGCAAC  
 CGCCCGTGTT CTTGCGGTAT CAACGAACGA ACGTTCTGAC ACCCCCGTTG  
 26151 ATCTCCTTCG CCCGCCGCTT TCTTCTCTAC CATCACGGCG TGGCCTTCCC  
 TAGAGGAAGC GGGCGGCGAA AGAAGAGATG GTAGTGCCGC ACCGGAAGGG  
 26201 CCGTAACATC CTGCATTACT ACCGTCATCT CTACAGCCCA TACTGCACCG  
 GGCATTGTAG GACGTAATGA TGGCAGTAGA GATGTCGGGT ATGACGTGGC  
 26251 GCGGCAGCGG CAGCAACAGC AGCGGCCACA CAGAAGCAAA GGCGACCGGA  
 CGCCGTCGCC GTCGTTGTG TCGCCGGTGT GTCTTCGTTT CCGCTGGCCT

FIG.9A-31

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26301 TAGCAAGACT CTGACAAAGC CCAAGAAATC CACAGCGGCG GCAGCAGCAG  
 ATCGTTCTGA GACTGTTTCG GGTCTTTAG GTGTCGCCGC CGTCGTCGTC

26351 GAGGAGGAGC GCTGCGTCTG GCGCCCAACG AACCCGTATC GACCCGCGAG  
 CTCCTCCTCG CGACGCAGAC CGCGGGTTGC TTGGGCATAG CTGGGCGCTC

26401 CTTAGAAACA GGATTTTTC CACTCTGTAT GCTATATTTT AACAGAGCAG  
 GAATCTTTGT CCTAAAAAGG GTGAGACATA CGATATAAAG TTGTCTCGTC

26451 GGGCCAAGAA CAAGAGCTGA AAATAAAAAA CAGGTCTCTG CGATCCCTCA  
 CCCGTTCTT GTTCTCGACT TTTATTTTTT GTCCAGAGAC GCTAGGGAGT

26501 CCCGCAGCTG CCTGTATCAC AAAAGCGAAG ATCAGCTTCG GCGCAGCTG  
 GGGCGTCGAC GGACATAGTG TTTTCGCTTC TAGTCGAAGC CGCGTGCGAC

26551 GAAGACGCGG AGGCTCTCTT CAGTAAATAC TGC GCGCTGA CTCTTAAGGA  
 CTTCTGCGCC TCCGAGAGAA GTCATTTATG ACGCGCGACT GAGAATTCCT

26601 CTAGTTTCGC GCCCTTCTC AAATTTAAGC GCGAAACTA CGTCATCTCC  
 GATCAAAGCG CGGGAAAGAG TTAAATTCG CGCTTTTGAT GCAGTAGAGG

26651 AGCGGCCACA CCCGGCGCCA GCACCTGTTG TCAGCGCCAT TATGAGCAAG  
 TCGCCGGTGT GGGCCGCGGT CGTGGACAAC AGTCGCGGTA ATACTCGTTC

26701 GAAATTCCCA CGCCCTACAT GTGGAGTTAC CAGCCACAAA TGGGACTTGC  
 CTTTAAGGGT GCGGGATGTA CACCTCAATG GTCGGTGTTT ACCCTGAACG

26751 GGCTGGAGCT GCCCAAGACT ACTCAACCCG AATAAACTAC ATGAGCGCGG  
 CCGACCTCGA CGGGTTCTGA TGAGTTGGGC TTATTTGATG TACTCGCGCC

26801 GACCCACAT GATATCCCGG GTCAACGGAA TACGCGCCCA CCGAAACCGA  
 CTGGGGTGTA CTATAGGGCC CAGTTGCCTT ATGCGCGGGT GGCTTTGGCT

26851 ATTCTCCTGG AACAGGCGGC TATTACCACC ACACCTCGTA ATAACCTTAA  
 TAAGAGGACC TTGTCCGCCG ATAATGGTGG TGTGGAGCAT TATTGGAATT

26901 TCCCCGTAGT TGGCCCGCTG CCCTGGTGTA CCAGGAAAGT CCCGCTCCCA  
 AGGGGCATCA ACCGGGCGAC GGGACCACAT GGTCTTTTCA GGGCGAGGGT

26951 CCACTGTGGT ACTTCCCAGA GACGCCCAGG CCGAAGTTCA GATGACTAAC  
 GGTGACACCA TGAAGGGTCT CTGCGGGTCC GGCTTCAAGT CTA CTGATTG

27001 TCAGGGGCGC AGCTTGCGGG CGGCTTTCGT CACAGGGTGC GGTGCGCCGG  
 AGTCCCCGCG TCGAACGCC GCGGAAAGCA GTGTCCACG CCAGCGGGCC

27051 GCAGGGTATA ACTCACCTGA CAATCAGAGG GCGAGGTATT CAGCTCAACG  
 CGTCCCATAT TGAGTGGACT GTTAGTCTCC CGCTCCATAA GTCGAGTTGC

27101 ACGAGTCGGT GAGCTCCTCG CTTGGTCTCC GTCCGGACGG GACATTTTCAG  
 TGCTCAGCCA CTCGAGGAGC GAACAGAGG CAGGCCTGCC CTGTAAAGTC

FIG.9A-32



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27151 ATCGGCGGCG CCGGCCGCTC TTCATTCACG CCTCGTCAGG CAATCCTAAC  
 TAGCCGCCGC GGCCGGCGAG AAGTAAGTGC GGAGCAGTCC GTTAGGATTG  
 27201 TCTGCAGACC TCGTCCTCTG AGCCGCGCTC TGGAGGCATT GGAAGTCTGC  
 AGACGTCTGG AGCAGGAGAC TCGGCGCGAG ACCTCCGTAA CCTTGAGACG  
 27251 AATTTATTGA GGAGTTTGTG CCATCGGTCT ACTTTAACCC CTTCTCGGGA  
 TTAAATAACT CCTCAAACAC GGTAGCCAGA TGAAATTGGG GAAGAGCCCT  
 27301 CCTCCCGGCC ACTATCCGGA TCAATTTATT CCTAACTTTG ACGCGGTAAA  
 GGAGGGCCGG TGATAGGCCCT AGTTAAATAA GGATTGAAAC TGCGCCATTT  
 27351 GGACTCGGCG GACGGCTACG ACTGAATGTT AAGTGGAGAG GCAGAGCAAC  
 CCTGAGCCGC CTGCCGATGC TGACTTACAA TTCACCTCTC CGTCTCGTTG  
 27401 TGCGCCTGAA ACACCTGGTC CACTGTCGCC GCCACAAGTG CTTTGCCCGC  
 ACGCGGACTT TGTGGACCAG GTGACAGCGG CGGTGTTTAC GAAACGGGCG  
 27451 GACTCCGGTG AGTTTTGCTA CTTTGAATTG CCCGAGGATC ATATCGAGGG  
 CTGAGGCCAC TCAAACGAT GAAACTTAAC GGGCTCCTAG TATAGCTCCC  
 27501 CCCGGCGCAC GCGTCCGGC TTACCGCCCA GGGAGAGCTT GCCCGTAGCC  
 GGGCCGCGTG CCGCAGGCCG AATGGCGGGT CCCTCTCGAA CGGGCATCGG  
 27551 TGATTGCGGA GTTTACCCAG CGCCCCCTGC TAGTTGAGCG GGACAGGGGA  
 ACTAAGCCCT CAAATGGGTC GCGGGGGACG ATCAACTCGC CCTGTCCCCT  
 27601 CCCTGTGTTT TCACTGTGAT TTGCAACTGT CCTAACCCCTG GATTACATCA  
 GGGACACAAG AGTGACACTA AACGTTGACA GGATTGGGAC CTAATGTAGT  
 27651 AGATCTTTGT TGCCATCTCT GTGCTGAGTA TAATAAATAC AGAAATTAAC  
 TCTAGAAACA ACGGTAGAGA CACGACTCAT ATTATTTATG TCTTTAATTT  
 27701 ATATACTGGG GCTCCTATCG CCATCCTGTA AACGCCACCG TCTTCACCCG  
 TATATGACCC CGAGGATAGC GGTAGGACAT TTGCGGTGGC AGAAGTGGGC  
 27751 CCCAAGCAAA CCAAGGCGAA CCTTACCTGG TACTTTTAAC ATCTCTCCCT  
 GGGTTCGTTT GGTTCCGCTT GGAATGGACC ATGAAAATTG TAGAGAGGGA  
 27801 CTGTGATTTA CAACAGTTTC AACCCAGACG GAGTGAGTCT ACGAGAGAAC  
 GACACTAAAT GTTGTCAAAG TTGGGTCTGC CTCACTCAGA TGCTCTCTTG  
 27851 CTCTCCGAGC TCAGCTACTC CATCAGAAAA AACACCACCC TCCTTACCTG  
 GAGAGGCTCG AGTCGATGAG GTAGTCTTTT TTGTGGTGGG AGGAATGGAC  
 27901 CCGGGAACGT ACGAGTGCCT CACCGGCCGC TGCACCACAC CTACCGCCTG  
 GGCCCTTGCA TGCTCACGCA GTGGCCGGCG ACGTGGTGTG GATGGCGGAC  
 27951 ACCGTAAACC AGACTTTTTTC CGGACAGACC TCAATAACTC TGTTTACCAG  
 TGGCATTTGG TCTGAAAAAG GCCTGTCTGG AGTTATTGAG ACAAATGGTC

FIG.9A-33

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28001 AACAGGAGGT GAGCTTAGAA AACCCTTAGG GTATTAGGCC AAAGGCGCAG  
 TTGTCTCCA CTCGAATCTT TTGGGAATCC CATAATCCGG TTTCCGCGTC  
 28051 CTA CTGTGGG GTTTATGAAC AATTCAAGCA ACTCTACGGG CTATTCTAAT  
 GATGACACCC CAAATACTTG TTAAGTTCGT TGAGATGCCC GATAAGATTA  
 28101 TCAGGTTTCT CTAGAATCGG GGTTGGGGTT ATTCTCTGTC TTGTGATTCT  
 AGTCCAAAGA GATCTTAGCC CCAACCCCAA TAAGAGACAG AACACTAAGA  
 28151 CTTTATTCTT ATACTAACGC TTCTCTGCCT AAGGCTCGCC GCCTGCTGTG  
 GAAATAAGAA TATGATTGCG AAGAGACGGA TTCCGAGCGG CGGACGACAC  
 28201 TGCACATTTG CATTTATTGT CAGCTTTTTA AACGCTGGGG TCGCCACCCA  
 ACGTGTA AAC GTAAATAACA GTCGAAAAAT TTGCGACCCC AGCGGTGGGT  
 28251 AGATGATTAG GTACATAATC CTAGGTTTAC TCACCCTTGC GTCAGCCAC  
 TCTACTAATC CATGTATTAG GATCCAAATG AGTGGGAACG CAGTCGGGTG  
 28301 GGTACCACCC AAAAGGTGGA TTTTAAGGAG CCAGCCTGTA ATGTTACATT  
 CCATGGTGGG TTTTCCACCT AAAATTCCTC GGTCGGACAT TACAATGTAA  
 28351 CGCAGCTGAA GCTAATGAGT GCACCACTCT TATAAAATGC ACCACAGAAC  
 CGGTCGACTT CGATTACTCA CGTGGTGAGA ATATTTTACG TGGTGTCTTG  
 28401 ATGAAAAGCT GCTTATTTCG CACAAAAACA AAATTGGCAA GTATGCTGTT  
 TACTTTTCGA CGAATAAGCG GTGTTTTTGT TTTAACC GTT CATACGACAA  
 28451 TATGCTATTT GGCAGCCAGG TGACACTACA GAGTATAATG TTACAGTTTT  
 ATACGATAAA CCGTCGGTCC ACTGTGATGT CTCATATTAC AATGTCAAAA  
 28501 CCAGGGTAAA AGTCATAAAA CTTTTATGTA TACTTTTCCA TTTTATGAAA  
 GGTCCCATTT TCAGTATTTT GAAAATACAT ATGAAAAGGT AAAATACTTT  
 28551 TGTGCGACAT TACCATGTAC ATGAGCAAAC AGTATAAGTT GTGGCCCCCA  
 ACACGCTGTA ATGGTACATG TACTCGTTTG TCATATTCAA CACCGGGGGT  
 28601 CAAAATTGTG TGGAAAACAC TGGCACTTTC TGCTGCACTG CTATGCTAAT  
 GTTTTAACAC ACCTTTTGTG ACCGTGAAAG ACGACGTGAC GATACGATTA  
 28651 TACAGTGCTC GCTTTGGTCT GTACCCTACT CTATATTAAA TACAAAAGCA  
 ATGTCACGAG CGAAACCAGA CATGGGATGA GATATAATTT ATGTTTTCTG  
 28701 GACGCAGCTT TATTGAGGAA AAGAAAATGC CTTAATTTAC TAAGTTACAA  
 CTGCGTCGAA ATA ACTCTT TTCTTTTACG GAATTAAATG ATTCAATGTT  
 28751 AGCTAATGTC ACCACTAACT GCTTTACTCG CTGCTTGCAA AACAAATTCA  
 TCGATTACAG TGGTGATTGA CGAAATGAGC GACGAACGTT TTGTTTAAAGT  
 28801 AAAAGTTAGC ATTATAATTA GAATAGGATT TAAACCCCCC GGTCAATTTCC  
 TTTTCAATCG TAATATTAAT CTTATCCTAA ATTTGGGGGG CCAAGTAAAGG

FIG.9A-34

43/70

28851 TGCTCAATAC CATTCCCCTG AACAAATTGAC TCTATGTGGG ATATGCTCCA  
ACGAGTTATG GTAAGGGGAC TTGTAACTG AGATACACCC TATACGAGGT

28901 GCGCTACAAC CTTGAAGTCA GGCTTCCTGG ATGTCAGCAT CTGACTTTGG  
CGCGATGTTG GAACTTCAGT CCGAAGGACC TACAGTCGTA GACTGAAACC

28951 CCAGCACCTG TCCCGCGGAT TTGTTCCAGT CCAACTACAG CGACCCACCC  
GGTCGTGGAC AGGGCGCCTA AACAAGGTCA GGTGATGTC GCTGGGTGGG

29001 TAACAGAGAT GACCAACACA ACCAACGCGG CCGCCGCTAC CGGACTTACA  
ATTGTCTCTA CTGGTTGTGT TGGTTGCGCC GCGGCGATG GCCTGAATGT

29051 TCTACCACAA ATACACCCCA AGTTTCTGCC TTTGTCAATA ACTGGGATAA  
AGATGGTGTT TATGTGGGGT TCAAAGACGG AAACAGTTAT TGACCCTATT

29101 CTTGGGCATG TGGTGGTTCT CCATAGCGCT TATGTTTGTG TGCCTTATTA  
GAACCCGTAC ACCACCAAGA GGTATCGCGA ATACAAACAT ACGGAATAAT

29151 TTATGTGGCT CATCTGCTGC CTAAAGCGCA AACGCGCCCG ACCACCCATC  
AATACACCGA GTAGACGACG GATTTCGCGT TTGCGCGGGC TGGTGGGTAG

29201 TATAGTCCCA TCATTGTGCT ACACCCAAAC AATGATGGAA TCCATAGATT  
ATATCAGGGT AGTAACACGA TGTGGGTTTG TTAACCTT AGGTATCTAA

29251 GGACGGACTG AAACACATGT TCTTTTCTCT TACAGTATGA TTAAATGAGA  
CCTGCCTGAC TTTGTGTACA AGAAAAGAGA ATGTCATACT AATTTACTCT

29301 CATGATTCTT CGAGTTTTTA TATTACTGAC CTTTGTGCG CTTTTTTGTG  
GTACTAAGGA GCTCAAAAAT ATAATGACTG GGAACAACGC GAAAAACAC

29351 CGTGCTCCAC ATTGGCTGCG GTTCTCACA TCGAAGTAGA CTGCATTCCA  
GCACGAGGTG TAACCGACGC CAAAGAGTGT AGCTTCATCT GACGTAAGGT

29401 GCCTTCACAG TCTATTTGCT TTACGGATTT GTCACCCTCA CGCTCATCTG  
CGGAAGTGTC AGATAAACGA AATGCCTAAA CAGTGGGAGT GCGAGTAGAC

29451 CAGCCTCATC ACTGTGGTCA TCGCCTTTAT CCAGTGCATT GACTGGGTCT  
GTCGGAGTAG TGACACCAGT AGCGGAAATA GGTCACGTAA CTGACCCAGA

29501 GTGTGCGCTT TGCATATCTC AGACACCATC CCCAGTACAG GGACAGGACT  
CACACGCGAA ACGTATAGAG TCTGTGGTAG GGGTCATGTC CCTGTCCTGA

29551 ATAGCTGAGC TTCTTAGAAT TCTTTAATTA TGAAATTTAC TGTGACTTTT  
TATCGACTCG AAGAATCTTA AGAAATTAAT ACTTTAAATG AACTGAAAA

29601 CTGCTGATTA TTTGCACCCT ATCTGCGTTT TGTTCCCGA CCTCCAAGCC  
GACGACTAAT AAACGTGGGA TAGACGCAA ACAAGGGGCT GGAGGTTCCG

29651 TCAAAGACAT ATATCATGCA GATTCACCTG TATATGGAAT ATTCCAAGTT  
AGTTTCTGTA TATAGTACGT CTAAGTGAGC ATATACCTTA TAAGGTTCAA

FIG.9A-35

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29701 GCTACAATGA AAAAAGCGAT CTTTCCGAAG CCTGGTTATA TGCAATCATC  
 CGATGTTACT TTTTTCGCTA GAAAGGCTTC GGACCAATAT ACGTTAGTAG  
 29751 TCTGTTATGG TGTTCTGCAG TACCATCTTA GCCCTAGCTA TATATCCCTA  
 AGACAATACC ACAAGACGTC ATGGTAGAAT CGGGATCGAT ATATAGGGAT  
 29801 CCTTGACATT GGCTGGAACG CAATAGATGC CATGAACCAC CCAACTTTCC  
 GGAACGTGTA CCGACCTTGC GTTATCTACG GTACTTGGTG GGTTGAAAGG  
 29851 CCGCGCCCGC TATGCTTCCA CTGCAACAAG TTGTTGCCGG CGGCTTTGTC  
 GCGCGGGGCG ATACGAAGGT GACGTTGTTC AACACGGCC GCCGAAACAG  
 29901 CCAGCCAATC AGCCTCGCCC ACCTTCTCCC ACCCCCACTG AAATCAGCTA  
 GGTGCGTTAG TCGGAGCGGG TGAAGAGGG TGGGGGTGAC TTTAGTCGAT  
 29951 CTTTAATCTA ACAGGAGGAG ATGACTGACA CCCTAGATCT AGAAATGGAC  
 GAAATTAGAT TGCCTCCTC TACTGACTGT GGGATCTAGA TCTTTACCTG  
 30001 GGAATTATTA CAGAGCAGCG CCTGCTAGAA AGACGCAGGG CAGCGGCCGA  
 CCTTAATAAT GTCTCGTCGC GGACGATCTT TCTGCGTCCC GTCGCCGGCT  
 30051 GCAACAGCGC ATGAATCAAG AGCTCCAAGA CATGGTTAAC TTGCACCAGT  
 CGTTGTCGCG TACTTAGTTC TCGAGGTTCT GTACCAATTG AACGTGGTCA  
 30101 GCAAAAGGGG TATCTTTTGT CTCGTAAAGC AGGCCAAAGT CACCTACGAC  
 CGTTTTCCCC ATAGAAAACA GAGCATTTCTG TCCGGTTTCA GTGGATGCTG  
 30151 AGTAATACCA CCGGACACCG CCTTAGCTAC AAGTTGCCAA CCAAGCGTCA  
 TCATTATGGT GGCCTGTGGC GGAATCGATG TTCAACGGTT GGTTTCGCAGT  
 30201 GAAATTGGTG GTCATGGTGG GAGAAAAGCC CATTACCATA ACTCAGCACT  
 CTTTAACCAC CAGTACCACC CTCTTTTCGG GTAATGGTAT TGAGTCGTGA  
 30251 CGGTAGAAAC CGAAGGCTGC ATTCACTCAC CTTGTCAAGG ACCTGAGGAT  
 GCCATCTTTG GCTTCCGACG TAAGTGAGTG GAACAGTTCC TGGACTCCTA  
 30301 CTCTGCACCC TTATTAAGAC CCTGTGCGGT CTCAAAGATC TTATTCCCTT  
 GAGACGTGGG AATAATTCTG GGACACGCCA GAGTTTCTAG AATAAGGGAA  
 30351 TAACTAATAA AAAAAAATAA TAAAGCATCA CTTACTTAAA ATCAGTTAGC  
 ATTGATTATT TTTTTTTATT ATTTCTGAGT GAATGAATTT TAGTCAATCG  
 30401 AAATTTCTGT CCAGTTTATT CAGCAGCACC TCCTTGCCCT CCTCCCAGCT  
 TTAAAGACA GGTCAAATAA GTCGTCGTGG AGGAACGGGA GGAGGGTCCA  
 30451 CTGGTATTGC AGCTTCCTCC TGGCTGCAAA CTTTCTCCAC AATCTAAATG  
 GACCATAACG TCGAAGGAGG ACCGACGTTT GAAAGAGGTG TTAGATTTAC  
 30501 GAATGTCAGT TTCCTCCTGT TCCTGTCCAT CCGCACCCAC TATCTTCATG  
 CTTACAGTCA AAGGAGGACA AGGACAGGTA GGCCTGGGTG ATAGAAGTAC

FIG.9A-36

45/70

30551 TTGTTGCAGA TGAAGCGCGC AAGACCGTCT GAAGATACCT TCAACCCCGT  
AACAAACGTCT ACTTCGCGCG TTCTGGCAGA CTTCTATGGA AGTTGGGGCA

30601 GTATCCATAT GACACGGAAA CCGGTCCTCC AACTGTGCCT TTTCTTACTC  
CATAGGTATA CTGTGCCTTT GGCCAGGAGG TTGACACGGA AAAGAATGAG

30651 CTCCCTTTGT ATCCCCAAT GGGTTTCAAG AGAGTCCCCC TGGGGTACTC  
GAGGGAAACA TAGGGGGTTA CCCAAAGTTC TCTCAGGGGG ACCCCATGAG

30701 TCTTTGCGCC TATCCGAACC TCTAGTTACC TCCAATGGCA TGCTTGCGCT  
AGAAACGCGG ATAGGCTTGG AGATCAATGG AGGTTACCGT ACGAACGCGA

30751 CAAAATGGGC AACGGCCTCT CTCTGGACGA GGCCGGCAAC CTTACCTCCC  
GTTTTACCCG TTGCCGGAGA GAGACCTGCT CCGGCCGTTG GAATGGAGGG

30801 AAAATGTAAC CACTGTGAGC CCACCTCTCA AAAAAACCAA GTCAAACATA  
TTTTACATTG GTGACACTCG GGTGGAGAGT TTTTTGGTT CAGTTTGTAT

30851 AACCTGGAAG TATCTGCACC CCTCACAGTT ACCTCAGAAG CCCTAACTGT  
TTGGACCTTT ATAGACGTGG GGAGTGTCAG TGGAGTCTTC GGGATTGACA

30901 GGCTGCCGCC GCACCTCTAA TGGTCGCGGG CAACACACTC ACCATGCAAT  
CCGACGGCGG CGTGGAGATT ACCAGCGCCC GTTGTGTGAG TGGTACGTTA

30951 CACAGGCCCC GCTAACCGTG CACGACTCCA AACTTAGCAT TGCCACCCAA  
GTGTCCGGGG CGATTGGCAC GTGCTGAGGT TTGAATCGTA ACGGTGGGTT

31001 GGACCCCTCA CAGTGTGAGA AGGAAAGCTA GCCCTGCAAA CATCAGGCCC  
CCTGGGGAGT GTCACAGTCT TCCTTTCGAT CGGGACGTTT GTAGTCCGGG

31051 CCTCACCACC ACCGATAGCA GTACCCTTAC TATCACTGCC TCACCCCTT  
GGAGTGGTGG TGGCTATCGT CATGGGAATG ATAGTGACGG AGTGGGGGAA

31101 TAACTACTGC CACTGGTAGC TTGGGCATTG ACTTGAAAGA GCCCATTTAT  
ATTGATGACG GTGACCATCG AACCCGTAAC TGAACCTTCT CGGGTAAATA

31151 ACACAAAATG GAAAACTAGG ACTAAAGTAC GGGGCTCCTT TGCATGTAAC  
TGTGTTTTAC CTTTTGATCC TGATTTTCATG CCCCAGGAA ACGTACATTG

31201 AGACGACCTA AACACTTTGA CCGTAGCAAC TGGTCCAGGT GTGACTATTA  
TCTGCTGGAT TTGTGAAACT GGCATCGTTG ACCAGGTCCA CACTGATAAT

31251 ATAATACTTC CTTGCAAAC AAAGTTACTG GAGCCTTGGG TTTTGATTCA  
TATTATGAAG GAACGTTTGA TTTCAATGAC CTCGGAACCC AAACTAAGT

31301 CAAGGCAATA TGCAACTTAA TGTAGCAGGA GGACTAAGGA TTGATTCTCA  
GTTCCGTTAT ACGTTGAATT ACATCGTCCT CCTGATTCTT AACTAAGAGT

31351 AAACAGACGC CTTATACTTG ATGTTAGTTA TCCGTTTGAT GCTCAAAACC  
TTTGTCTGCG GAATATGAAC TACAATCAAT AGGCAAACTA CGAGTTTGG

FIG.9A-37

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31401 AACTAAATCT AAGACTAGGA CAGGGCCCTC TTTTATAAA CTCAGCCCAC  
TTGATTTAGA TTCTGATCCT GTCCCGGGAG AAAAATATTT GAGTCGGGTG

31451 AACTTGGATA TTAACACAA CAAAGGCCTT TACTTGTTTA CAGCTTCAAA  
TTGAACCTAT AATTGATGTT GTTCCGGAA ATGAACAAAT GTCGAAGTTT

31501 CAATTCCAAA AAGCTTGAGG TTAACCTAAG CACTGCCAAG GGGTTGATGT  
GTTAAGGTTT TTCGAACTCC AATTGGATTC GTGACGGTTC CCCAACTACA

31551 TTGACGCTAC AGCCATAGCC ATTAATGCAG GAGATGGGCT TGAATTTGGT  
AACTGCGATG TCGGTATCGG TAATTACGTC CTCTACCCGA ACTTAAACCA

31601 TCACCTAATG CACCAAACAC AAATCCCCTC AAAACAAAAA TTGGCCATGG  
AGTGGATTAC GTGGTTTGTG TTTAGGGGAG TTTTGTTTTT AACCGGTACC

31651 CCTAGAATTT GATTCAAACA AGGCTATGGT TCCTAAACTA GGAAGTGGCC  
GGATCTTAAA CTAAGTTTGT TCCGATACCA AGGATTTGAT CCTTGACCGG

31701 TTAGTTTTGA CAGCACAGGT GCCATTACAG TAGGAAACAA AAATAATGAT  
AATCAAAACT GTCGTGTCCA CGGTAATGTC ATCCTTTGTT TTTATTACTA

31751 AAGCTAACTT TGTGGACCAC ACCAGCTCCA TCTCCTAACT GTAGACTAAA  
TTCGATTGAA ACACCTGGTG TGGTCGAGGT AGAGGATTGA CATCTGATTT

31801 TGCAGAGAAA GATGCTAAAC TCACTTTGGT CTTAACAAAA TGTGGCAGTC  
ACGTCTCTTT CTACGATTTG AGTGAAACCA GAATTGTTTT ACACCGTCAG

31851 AAATACTTGC TACAGTTTCA GTTTTGGCTG TTAAAGGCAG TTTGGCTCCA  
TTTATGAACG ATGTCAAAGT CAAAACCGAC AATTTCCGTC AAACCGAGGT

31901 ATATCTGGAA CAGTTCAAAG TGCTCATCTT ATTATAAGAT TTGACGAAAA  
TATAGACCTT GTCAAGTTTC ACGAGTAGAA TAATATTCTA AACTGCTTTT

31951 TGGAGTGCTA CTAAACAATT CCTTCCTGGA CCCAGAATAT TGGAACCTTTA  
ACCTCACGAT GATTTGTTAA GGAAGGACCT GGGTCTTATA ACCTTGAAAT

32001 GAAATGGAGA TCTTACTGAA GGCACAGCCT ATACAAACGC TGTTGGATTT  
CTTTACCTCT AGAATGACTT CCGTGTCGGA TATGTTTGCG ACAACCTAAA

32051 ATGCCTAACC TATCAGCTTA TCCAAAATCT CACGGTAAAA CTGCCAAAAG  
TACGGATTGG ATAGTCGAAT AGGTTTTAGA GTGCCATTTT GACGGTTTTT

32101 TAACATTGTC AGTCAAGTTT ACTTAAACGG AGACAAAACCT AAACCTGTAA  
ATTGTAACAG TCAGTTCAAA TGAATTTGCC TCTGTTTTGA TTTGGACATT

32151 CACTAACCAT TACACTAAAC GGTACACAGG AAACAGGAGA CACAACCTCA  
GTGATTGGTA ATGTGATTTG CCATGTGTCC TTTGTCTCT GTGTTGAGGT

32201 AGTGCATACT CTATGTCATT TTCATGGGAC TGGTCTGGCC ACAACTACAT  
TCACGTATGA GATACAGTAA AAGTACCCTG ACCAGACCGG TGTGATGTA

FIG.9A-38

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|       |            |            |             |            |            |
|-------|------------|------------|-------------|------------|------------|
| 32251 | TAATGAAATA | TTTGCCACAT | CCTCTTACAC  | TTTTTCATAC | ATTGCCCAAG |
|       | ATTACTTTAT | AAACGGTGTA | GGAGAATGTG  | AAAAAGTATG | TAACGGGTTC |
| 32301 | AATAAAGAAT | CGTTTGTGTT | ATGTTTCAAC  | GTGTTTATTT | TTCAATTGCA |
|       | TTATTTCTTA | GCAAACACAA | TACAAAGTTG  | CACAAATAAA | AAGTTAACGT |
| 32351 | GAAAATTTCA | AGTCATTTTT | CATTCAAGTAG | TATAGCCCCA | CCACCACATA |
|       | CTTTTAAAGT | TCAGTAAAAA | GTAAGTCATC  | ATATCGGGGT | GGTGGTGTAT |
| 32401 | GCTTATACAG | ATCACCGTAC | CTTAATCAAA  | CTCACAGAAC | CCTAGTATTC |
|       | CGAATATGTC | TAGTGGCATG | GAATTAGTTT  | GAGTGTCTTG | GGATCATAAG |
| 32451 | AACCTGCCAC | CTCCCTCCCA | ACACACAGAG  | TACACAGTCC | TTTCTCCCCG |
|       | TTGGACGGTG | GAGGGAGGGT | TGTGTGTCTC  | ATGTGTCAGG | AAAGAGGGGC |
| 32501 | GCTGGCCTTA | AAAAGCATCA | TATCATGGGT  | AACAGACATA | TTCTTAGGTG |
|       | CGACCGGAAT | TTTTCGTAGT | ATAGTACCCA  | TTGTCTGTAT | AAGAATCCAC |
| 32551 | TTATATTCCA | CACGGTTTCC | TGTCGAGCCA  | AACGCTCATC | AGTGATATTA |
|       | AATATAAGGT | GTGCCAAAGG | ACAGCTCGGT  | TTGCGAGTAG | TCACTATAAT |
| 32601 | ATAAACTCCC | CGGGCAGCTC | ACTTAAGTTC  | ATGTCGCTGT | CCAGCTGCTG |
|       | TATTTGAGGG | GCCCGTCGAG | TGAATTCAAG  | TACAGCGACA | GGTCGACGAC |
| 32651 | AGCCACAGGC | TGCTGTCCAA | CTTGCGGTTG  | CTTAACGGGC | GGCGAAGGAG |
|       | TCGGTGTCCG | ACGACAGGTT | GAACGCCAAC  | GAATTGCCCC | CCGCTTCCTC |
| 32701 | AAGTCCACGC | CTACATGGGG | GTAGAGTCAT  | AATCGTGCAT | CAGGATAGGG |
|       | TTCAGGTGCG | GATGTACCCC | CATCTCAGTA  | TTAGCACGTA | GTCCTATCCC |
| 32751 | CGGTGGTGCT | GCAGCAGCGC | GCGAATAAAC  | TGCTGCCGCC | GCCGCTCCGT |
|       | GCCACCACGA | CGTCGTCGCG | CGCTTATTTG  | ACGACGGCGG | CGGCGAGGCA |
| 32801 | CCTGCAGGAA | TACAACATGG | CAGTGGTCTC  | CTCAGCGATG | ATTCGCACCG |
|       | GGACGTCCTT | ATGTTGTACC | GTCACCAGAG  | GAGTCGCTAC | TAAGCGTGGC |
| 32851 | CCCGCAGCAT | AAGGCGCCTT | GTCCTCCGGG  | CACAGCAGCG | CACCCTGATC |
|       | GGGCGTCGTA | TTCCGCGGAA | CAGGAGGCCC  | GTGTCGTCGC | GTGGGACTAG |
| 32901 | TCACTTAAAT | CAGCACAGTA | ACTGCAGCAC  | AGCACCACAA | TATTGTTCAA |
|       | AGTGAATTTA | GTCGTGTCAT | TGACGTCGTG  | TCGTGGTGTT | ATAACAAGTT |
| 32951 | AATCCCACAG | TGCAAGGCGC | TGTATCCAAA  | GCTCATGGCG | GGGACCACAG |
|       | TTAGGGTGTC | ACGTTCCGCG | ACATAGGTTT  | CGAGTACCGC | CCCTGGTGTC |
| 33001 | AACCCACGTG | GCCATCATAC | CACAAGCGCA  | GGTAGATTAA | GTGGCGACCC |
|       | TTGGGTGCAC | CGGTAGTATG | GTGTTGCGGT  | CCATCTAATT | CACCGCTGGG |
| 33051 | CTCATAAACA | CGCTGGACAT | AAACATTACC  | TCTTTTGGCA | TGTTGTAATT |
|       | GAGTATTTGT | GCGACCTGTA | TTTGTAATGG  | AGAAAACCGT | ACAACATTAA |

FIG.9A-39

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33101 CACCACCTCC CGGTACCATA TAAACCTCTG ATTAACATG GCGCCATCCA  
 GTGGTGGAGG GCCATGGTAT ATTTGGAGAC TAATTTGTAC CGCGGTAGGT  
 33151 CCACCATCCT AAACCAGCTG GCCAAAACCT GCCCGCCGGC TATACACTGC  
 GGTGGTAGGA TTTGGTCGAC CGGTTTTGGA CGGGCGGCCG ATATGTGACG  
 33201 AGGGAACCGG GACTGGAACA ATGACAGTGG AGAGCCCAGG ACTCGTAACC  
 TCCCTTGGCC CTGACCTTGT TACTGTCACC TCTCGGTCC TGAGCATTGG  
 33251 ATGGATCATC ATGCTCGTCA TGATATCAAT GTTGGCACA CACAGGCACA  
 TACCTAGTAG TACGAGCAGT ACTATAGTTA CAACCGTGTT GTGTCCGTGT  
 33301 CGTGCATACA CTTCTCAGG ATTACAAGCT CCTCCCGCGT TAGAACCATA  
 GCACGTATGT GAAGGAGTCC TAATGTTTGA GGAGGGCGCA ATCTTGGTAT  
 33351 TCCCAGGGAA CAACCCATTG CTGAATCAGC GTAAATCCCA CACTGCAGGG  
 AGGGTCCCTT GTTGGGTAAG GACTTAGTCG CATTTAGGGT GTGACGTCCC  
 33401 AAGACCTCGC ACGTAACTCA CGTTGTGCAT TGTCAAAGTG TTACATTCGG  
 TTCTGGAGCG TGCATTGAGT GCAACACGTA ACAGTTTCAC AATGTAAGCC  
 33451 GCAGCAGCGG ATGATCCTCC AGTATGGTAG CGCGGGTTTC TGTCTCAAAA  
 CGTCGTGCGC TACTAGGAGG TCATACCATC GCGCCCAAAG ACAGAGTTTT  
 33501 GGAGGTAGAC GATCCCTACT GTACGGAGTG CGCCGAGACA ACCGAGATCG  
 CCTCCATCTG CTAGGGATGA CATGCCTCAC GCGGCTCTGT TGGCTCTAGC  
 33551 TGTTGGTCGT AGTGTGATGC CAAATGGAAC GCCGGACGTA GTCATATTTT  
 ACAACCAGCA TCACAGTACG GTTTACCTTG CGGCCTGCAT CAGTATAAAG  
 33601 CTGAAGCAAA ACCAGGTGCG GGCCTGACAA ACAGATCTGC GTCTCCGGTC  
 GACTTCGTTT TGGTCCACGC CCGCACTGTT TGTCTAGACG CAGAGGCCAG  
 33651 TCGCCGCTTA GATCGCTCTG TGTAGTAGTT GTAGTATATC CACTCTCTCA  
 AGCGGCGAAT CTAGCGAGAC ACATCATCAA CATCATATAG GTGAGAGAGT  
 33701 AAGCATCCAG GCGCCCCCTG GCTTCGGGTT CTATGTAAAC TCCTTCATGC  
 TTCGTAGGTC CGCGGGGGAC CGAAGCCCAA GATACATTTG AGGAAGTACG  
 33751 GCCGCTGCCC TGATAACATC CACCACCGCA GAATAAGCCA CACCCAGCCA  
 CGGCGACGGG ACTATTGTAG GTGGTGGCGT CTTATTCGGT GTGGGTGCGT  
 33801 ACCTACACAT TCGTTCTGCG AGTCACACAC GGGAGGAGCG GGAAGAGCTG  
 TGGATGTGTA AGCAAGACGC TCAGTGTGTG CCCTCCTCGC CCTTCTCGAC  
 33851 GAAGAACCAT GTTTTTTTTT TTATTCCAAA AGATTATCCA AAACCTCAAA  
 CTTCTTGGTA CAAAAAATAA AATAAGGTTT TCTAATAGGT TTTGGAGTTT  
 33901 ATGAAGATCT ATTAAGTGAA CGCGCTCCCC TCCGGTGGCG TGGTCAAAC  
 TACTTCTAGA TAATTCACCT GCGCGAGGGG AGGCCACCGC ACCAGTTTGA

FIG.9A-40



49/70

33951 CTACAGCCAA AGAACAGATA ATGGCATTG TAAGATGTTG CACAATGGCT  
 GATGTCGGT TCTGTCTAT TACCGTAAAC ATTCTACAAC GTGTTACCGA  
 34001 TCCAAAAGGC AAACGGCCCT CACGTCCAAG TGGACGTAAA GGCTAAACCC  
 AGGTTTTCCG TTGCGCGGA GTGCAGGTC ACCTGCATT CCGATTGGG  
 34051 TTCAGGGTGA ATCTCCTCTA TAAACATTCC AGCACCTTCA ACCATGCCCA  
 AAGTCCCACT TAGAGGAGAT ATTTGTAAGG TCGTGGAAGT TGGTACGGGT  
 34101 AATAATTCTC ATCTCGCCAC CTTCTCAATA TATCTCTAAG CAAATCCCGA  
 TTATTAAGAG TAGAGCGGTG GAAGAGTTAT ATAGAGATTC GTTTAGGGCT  
 34151 ATATTAAGTC CGGCCATTGT AAAAATCTGC TCCAGAGCGC CCTCCACCTT  
 TATAATTGAG GCCGGTAACA TTTTAGACG AGGTCTCGCG GGAGGTGGAA  
 34201 CAGCCTCAAG CAGCGAATCA TGATTGCAAA AATTCAGGTT CCTCACAGAC  
 GTCGGAGTTC GTCGCTTAGT ACTAACGTTT TTAAGTCCAA GGAGTGTCTG  
 34251 CTGTATAAGA TTCAAAAGCG GAACATTAAC AAAAATACCG CGATCCCGTA  
 GACATATTCT AAGTTTTCGC CTTGTAATTG TTTTATGGC GCTAGGGCAT  
 34301 GGTCCCTTCG CAGGGCCAGC TGAACATAAT CGTGCAGGTC TGCACGGACC  
 CCAGGGAAGC GTCCCGGTG ACTTGTTATTA GCACGTCCAG ACGTGCCTGG  
 34351 AGCGCGGCCA CTTCCCCGCC AGGAACCATG ACAAAGAAGC CCACACTGAT  
 TCGCGCCGGT GAAGGGGCGG TCCTTGGTAC TGTTTTCTTG GGTGTGACTA  
 34401 TATGACACGC ATACTCGGAG CTATGCTAAC CAGCGTAGCC CCGATGTAAG  
 ATACTGTGCG TATGAGCCTC GATACGATTG GTCGCATCGG GGCTACATTC  
 34451 CTTGTTGCAT GGGCGGCGAT ATAAATGCA AGGTGCTGCT CAAAAAATCA  
 GAACAACGTA CCCGCCGCTA TATTTTACGT TCCACGACGA GTTTTTTAGT  
 34501 GGCAAAGCCT CGCGCAAAAA AGAAAGCACA TCGTAGTCAT GCTCATGCAG  
 CCGTTTCGGA GCGCGTTTTT TCTTTCGTGT AGCATCAGTA CGAGTACGTC  
 34551 ATAAAGGCAG GTAAGCTCCG GAACCACCAC AGAAAAAGAC ACCATTTTTC  
 TATTTCCGTC CATTGAGGC CTTGGTGGTG TCTTTTCTG TGGTAAAAAG  
 34601 TCTCAAACAT GTCTGCGGGT TTCTGCATAA ACACAAAATA AAATAACAAA  
 AGAGTTTGTA CAGACGCCCA AAGACGTATT TGTGTTTTAT TTTATTGTTT  
 34651 AAAACATTTA AACATTAGAA GCCTGTCTTA CAACAGGAAA AACAACCCTT  
 TTTTGTAAT TTGTAATCTT CGGACAGAAT GTTGTCCTTT TTGTTGGGAA  
 34701 ATAAGCATAA GACGGACTAC GGCCATGCCG GCGTGACCGT AAAAAAATG  
 TATTCGTATT CTGCCTGATG CCGGTACGGC CGCACTGGCA TTTTTTGAC  
 34751 GTCACCGTGA TTA AAAAGCA CCACCGACAG CTCCTCGGTC ATGTCCGGAG  
 CAGTGGCACT AATTTTTCGT GGTGGCTGTC GAGGAGCCAG TACAGGCCTC

FIG.9A-41

50/70

34801 TCATAATGTA AGACTCGGTA AACACATCAG GTTGATTCAC ATCGGTCAGT  
 AGTATTACAT TCTGAGCCAT TTGTGTAGTC CAACTAAGTG TAGCCAGTCA  
 34851 GCTAAAAAGC GACCGAAATA GCCCGGGGGA ATACATACCC GCAGGCGTAG  
 CGATTTTTCG CTGGCTTTAT CGGGCCCCCT TATGTATGGG CGTCCGCATC  
 34901 AGACAACATT ACAGCCCCCA TAGGAGGTAT AACAAAATTA ATAGGAGAGA  
 TCTGTTGTAA TGTCGGGGGT ATCCTCCATA TTGTTTAAAT TATCCTCTCT  
 34951 AAAACACATA AACACCTGAA AAACCCTCCT GCCTAGGCAA AATAGCACCC  
 TTTTGTGTAT TTGTGGACTT TTTGGGAGGA CGGATCCGTT TTATCGTGGG  
 35001 TCCCCTCCA GAACAACATA CAGCGCTTCC ACAGCGGCAG CCATAACAGT  
 AGGGCGAGGT CTTGTTGTAT GTCGCGAAGG TGTCGCCGTC GGTATTGTCA  
 35051 CAGCCTTACC AGTAAAAAAG AAAACCTATT AAAAAAACAC CACTCGACAC  
 GTCGGAATGG TCATTTTTTC TTTTGATAA TTTTTTGTG GTGAGCTGTG  
 35101 GGCACCAGCT CAATCAGTCA CAGTGTAATA AAGGGCCAAG TGCAGAGCGA  
 CCGTGGTCTGA GTTAGTCAGT GTCACATTTT TTCCCGGTTT ACGTCTCGCT  
 35151 GTATATATAG GACTAAAAAA TGACGTAACG GTTAAAGTCC ACAAAAAACA  
 CATATATATC CTGATTTTTT ACTGCATTGC CAATTTTCAGG TGTTTTTTGT  
 35201 CCCAGAAAAC CGCACGCGAA CCTACGCCCA GAAACGAAAG CCAAAAAACC  
 GGGTCTTTTG GCGTGCGCTT GGATGCGGGT CTTTGCTTTC GGTTTTTTGG  
 35251 CACAATTCC TCAAATCGTC ACTTCCGTTT TCCCACGTTA CGTCACTTCC  
 GTGTTGAAGG AGTTTAGCAG TGAAGGCAAA AGGGTGCAAT GCAGTGAAGG  
 35301 CATTTTAAGA AACTACAAT TCCCAACACA TACAAGTTAC TCCGCCCTAA  
 GTAAAATTCT TTTGATGTGA AGGGTTGTGT ATGTTCAATG AGGCGGGATT  
 35351 AACCTACGTC ACCCGCCCCG TTCCACGCC CCGCGCCACG TCACAACTC  
 TTGGATGCAG TGGCGGGGCG AAGGGTGCGG GGC GCGGTGC AGTGTTTGAG  
 35401 CACCCCTCA TTATCATATT GGCTTCAATC CAAAATAAGG TATATTATTG  
 GTGGGGGAGT AATAGTATAA CCGAAGTTAG GTTTTATTCC ATATAATAAC  
 PacI  
 35451 ATGATGTTAA TTAAGAATTC GGATCTGCGA CGCGAGGCTG GATGGCCTTC  
 TACTACAATT AATCTTAAG CCTAGACGCT GCGCTCCGAC CTACCGGAAG  
 35501 CCCATTATGA TTCTTCTCGC TTCCGGCGGC ATCGGGATGC CCGCGTTGCA  
 GGGTAATACT AAGAAGAGCG AAGGCCGCCG TAGCCCTACG GGC GCAACGT  
 35551 GGCCATGCTG TCCAGGCAGG TAGATGACGA CCATCAGGGA CAGCTTCAAG  
 CCGGTACGAC AGGTCCGTCC ATCTACTGCT GGTAGTCCCT GTCGAAGTTC

FIG.9A-42

51/70

35601 GCCAGCAAAA GGCCAGGAAC CGTAAAAAGG CCGCGTTGCT GGCGTTTTTC  
 CGGTCGTTTT CCGGTCCTTG GCATTTTTCC GGCGCAACGA CCGCAAAAAG  
 35651 CATAGGCTCC GCCCCCTGA CGAGCATCAC AAAAATCGAC GCTCAAGTCA  
 GTATCCGAGG CGGGGGGACT GCTCGTAGTG TTTTAGCTG CGAGTTCAGT  
 35701 GAGGTGGCGA AACCCGACAG GACTATAAAG ATACCAGGCG TTTCCCCTG  
 CTCCACCGCT TTGGGCTGTC CTGATATTC TATGGTCCGC AAAGGGGGAC  
 35751 GAAGCTCCCT CGTGCGCTCT CCTGTTCCGA CCCTGCCGCT TACCGGATAC  
 CTTGAGAGGA GCACGCGAGA GGACAAGGCT GGGACGGCGA ATGGCCTATG  
 35801 CTGTCCGCCT TTCTCCCTTC GGAAGCGTG GCGCTTTCTC ATAGCTCACG  
 GACAGGCGGA AAGAGGGAAG CCCTTCGCAC CGCGAAAGAG TATCGAGTGC  
 35851 CTGTAGGTAT CTCAGTTCGG TGTAGGTCGT TCGCTCCAAG CTGGGCTGTG  
 GACATCCATA GAGTCAAGCC ACATCCAGCA AGCGAGGTTC GACCCGACAC  
 35901 TGCACGAACC CCCCGTTTCA CCCGACCGCT GCGCCTTATC CGGTAACAT  
 ACGTGCTTGG GGGGCAAGTC GGGCTGGCGA CGCGGAATAG GCCATTGATA  
 35951 CGTCTTGAGT CCAACCCGGT AAGACACGAC TTATCGCCAC TGGCAGCAGC  
 GCAGAACTCA GGTTGGGCCA TTCTGTGCTG AATAGCGGTG ACCGTCGTG  
 36001 CACTGGTAAC AGGATTAGCA GAGCGAGGTA TGTAGGCGGT GCTACAGAGT  
 GTGACCATTG TCCTAATCGT CTCGCTCCAT ACATCCGCCA CGATGTCTCA  
 36051 TCTTGAAGTG GTGGCCTAAC TACGGCTACA CTAGAAGGAC AGTATTTGGT  
 AGAACTTCAC CACCGGATTG ATGCCGATGT GATCTTCCTG TCATAAACCA  
 36101 ATCTGCGCTC TGCTGAAGCC AGTTACCTTC GGAAAAAGAG TTGGTAGCTC  
 TAGACGCGAG ACGACTTCGG TCAATGGAAG CCTTTTTCTC AACCATCGAG  
 36151 TTGATCCGGC AAACAAACCA CCGCTGGTAG CCGTGGTTTT TTTGTTTGCA  
 AACTAGGCCG TTTGTTTGGT GGCGACCATC GCCACCAAAA AAACAAACGT  
 36201 AGCAGCAGAT TACGCGCAGA AAAAAAGGAT CTCAAGAAGA TCCTTTGATC  
 TCGTCGTCTA ATGCGCGTCT TTTTTCTA GAGTTCTTCT AGGAACTAG  
 36251 TTTTCTACGG GGTCTGACGC TCAGTGGAAC GAAAACTCAC GTTAAGGGAT  
 AAAAGATGCC CCAGACTGCG AGTCACCTTG CTTTGTAGTG CAATCCCTA  
 36301 TTTGGTCATG AGATTATCAA AAAGGATCTT CACCTAGATC CTTTTAAATC  
 AAACCAGTAC TCTAATAGTT TTTCTAGAA GTGGATCTAG GAAAATTTAG  
 36351 AATCTAAAGT ATATATGAGT AAAGTTGGTC TGACAGTTAC CAATGCTTAA  
 TTAGATTTCA TATATACTCA TTTGAACCAG ACTGTCAATG GTTACGAATT  
 36401 TCAGTGAGGC ACCTATCTCA GCGATCTGTC TATTTGTTTC ATCCATAGTT  
 AGTCACTCCG TGGATAGAGT CGCTAGACAG ATAAAGCAAG TAGGTATCAA

FIG.9A-43

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36451 GCCTGACTCC CCGTCGTGTA GATAACTACG ATACGGGAGG GCTTACCATC  
 CGGACTGAGG GGCAGCACAT CTATTGATGC TATGCCCTCC CGAATGGTAG  
 36501 TGGCCCCAGT GCTGCAATGA TACCGCGAGA CCCACGCTCA CCGGCTCCAG  
 ACCGGGGTCA CGACGTTACT ATGGCGCTCT GGGTGCGAGT GGCCGAGGTC  
 36551 ATTTATCAGC AATAAACCAG CCAGCCGGAA GGGCCGAGCG CAGAAGTGGT  
 TAAATAGTCG TTATTTGGTC GGTGCGCCTT CCCGGCTCGC GTCTTCACCA  
 36601 CCTGCAACTT TATCCGCCTC CATCCAGTCT ATTAATTGTT GCCGGGAAGC  
 GGACGTTGAA ATAGGCGGAG GTAGGTCAGA TAATTAACAA CGGCCCTTCG  
 36651 TAGAGTAAGT AGTTCGCCAG TTAATAGTTT GCGCAACGTT GTTGCCATTG  
 ATCTCATTCA TCAAGCGGTC AATTATCAAA CGCGTTGCAA CAACGGTAAC  
 36701 CTACAGGCAT CGTGGTGTCA CGCTCGTCGT TTGGTATGGC TTCATTACAGC  
 GATGTCCGTA GCACCACAGT GCGAGCAGCA AACCATAACG AAGTAAGTCG  
 36751 TCCGGTTCCC AACGATCAAG GCGAGTTACA TGATCCCCCA TGTTGTGCAA  
 AGGCCAAGGG TTGCTAGTTC CGCTCAATGT ACTAGGGGGT ACAACACGTT  
 36801 AAAAGCGGTT AGCTCCTTCG GTCCTCCGAT CGTTGTCAGA AGTAAGTTGG  
 TTTTCGCCAA TCGAGGAAGC CAGGAGGCTA GCAACAGTCT TCATTCAACC  
 36851 CCGCAGTGTT ATCACTCATG GTTATGGCAG CACTGCATAA TTCTCTTACT  
 GGCGTCACAA TAGTGAGTAC CAATACCGTC GTGACGTATT AAGAGAATGA  
 36901 GTCATGCCAT CCGTAAGATG CTTTTCTGTG ACTGGTGAGT ACTCAACCAA  
 CAGTACGGTA GGCATTCTAC GAAAAGACAC TGACCACTCA TGAGTTGGTT  
 36951 GTCATTCTGA GAATAGTGTA TGC GGCGACC GAGTTGCTCT TGCCCGGCGT  
 CAGTAAGACT CTTATCACAT ACGCCGCTGG CTCAACGAGA ACGGGCCGCA  
 37001 CAACACGGGA TAATACCGCG CCACATAGCA GAACTTTAAA AGTGCTCATC  
 GTTGTGCCCT ATTATGGCGC GGTGTATCGT CTTGAAATTT TCACGAGTAG  
 37051 ATTGGA AAAC GTTCTTCGGG GCGAAAAC TC AAGGATCT TACCGCTGTT  
 TAACCTTTTG CAAGAAGCCC CGCTTTTGAG AGTTCCTAGA ATGGCGACAA  
 37101 GAGATCCAGT TCGATGTAAC CCACTCGTGC ACCCAACTGA TCTTCAGCAT  
 CTCTAGGTCA AGCTACATTG GGTGAGCACG TGGGTTGACT AGAAGTCGTA  
 37151 CTTTTACTTT CACCAGCGTT TCTGGGTGAG CAAAAACAGG AAGGCAAAAT  
 GAAAATGAAA GTGGTCGCAA AGACCCACTC GTTTTTGTCC TTCCGTTTTA  
 37201 GCCGCAAAAA AGGGAATAAG GGCGACACGG AAATGTTGAA TACTCATACT  
 CGGCGTTTTT TCCCTTATTC CCGCTGTGCC TTTACAACTT ATGAGTATGA  
 37251 CTTCTTTTTT CAATATTATT GAAGCATTTA TCAGGGTTAT TGTCTCATGA  
 GAAGGAAAAA GTTATAATAA CTTGTAAT AGTCCCAATA ACAGAGTACT

FIG.9A-44

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37301 GCGGATACAT ATTTGAATGT ATTTAGAAAA ATAAACAAAT AGGGGTTCCG  
CGCCTATGTA TAAACTTACA TAAATCTTTT TATTTGTTTA TCCCAAGGC

37351 CGCACATTTT CCGGAAAAGT GCCACCTGAC GTCTAAGAAA CCATTATTAT  
GCGTGTAAG GGGCTTTTCA CGGTGGACTG CAGATTCTTT GGTAATAATA

37401 CATGACATTA ACCTATAAAA ATAGGCGTAT CACGAGGCC TTTTCGTCTTC  
GTA CTGTAAT TGGATATTTT TATCCGCATA GTGCTCCGGG AAAGCAGAAG

37451 AAGAATTGGA TCCGAATTCT TAAT  
TTCTTAACCT AGGCTTAAGA ATTA

FIG.9A-45

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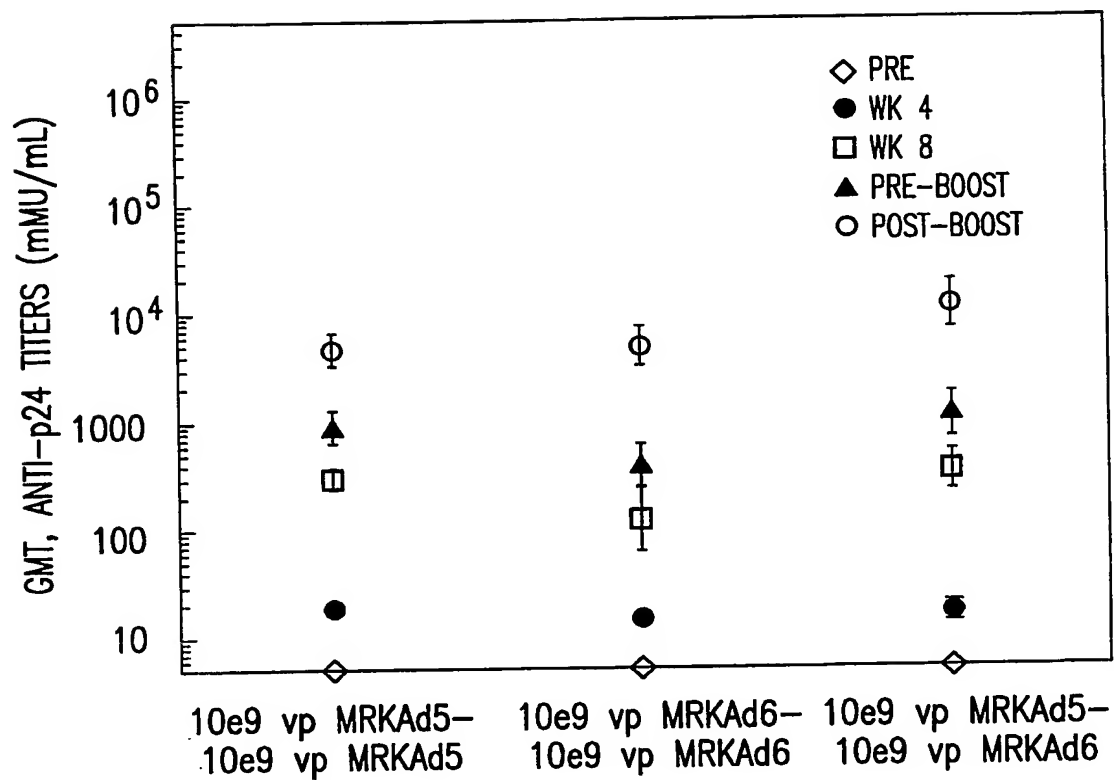


FIG.10

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1 CATCATCAAT AATATACCTT ATTTTGGATT GAAGCCAATA TGATAATGAG GGGGTGGAGT
61 TTGTGACGTG GCGCGGGGCG TGGGAACGGG GCGGGTGACG TAGTAGTGTG GCGGAAGTGT
121 GATGTTGTAA GTGTGGCGGA ACACATGTAA GCGCCGGATG TGGTAAAAGT GACGTTTTTG
181 GTGTGCGCCG GTGTACACGG GAAGTGACAA TTTTCGCGCG GTTTTAGGCG GATGTTGTAG
241 TAAATTTGGG CGTAACCAAG TAATATTTGG CCATTTTCGC GGGAAAACGT AATAAGAGGA
301 AGTGAAATCT GAATAATTCT GTGTACTCA TAGCGGTAA TATTTGTCTA GGGCCGCGGG
361 GACTTTGACC GTTTACGTGG AGACTCGCCC AGGTGTTTT CTCAGGTGT TTCCGCGTTC
421 CGGGTCAAAG TTGGCGTTTT ATTATTATAG TCAGCTGACG CGCAGTGTAT TTATACCCGG
481 TGAGTTCCTC AAGAGGCCAC TCTTGAGTGC CAGCGAGTAG AGTTTTCTCC TCCGAGCCGC
541 TCCGACACCG GGAAGTAAAA TGAGACATAT TATCTGCCAC GGAGGTGTTA TTACCGAAGA
601 AATGGCCGCC AGTCTTTTGG ACCAGCTGAT CGAAGAGGTA CTGGCTGATA ATCTTCCACC
661 TCCTAGCCAT TTTGAACCAC CTACCCTTCA CGAACTGTAT GATTTAGACG TGACGGCCCC
721 CGAAGATCCC AACGAGGAGG CGGTTTCGCA GATTTTTCCT GAGTCTGTAA TGTTGGCGGT
781 GCAGGAAGGG ATTGACTTAT TCACTTTTCC GCCGGCGCCC GGTTCTCCGG AGCCGCCTCA
841 CCTTTCCTCG CAGCCCGAGC AGCCGGAGCA GAGAGCCTTG GGTCCGGTTT CTATGCCAAA
901 CCTGTGCGCG GAGGTGATCG ATCTTACCTG CCACGAGGCT GGCTTTCAC CCAGTGACGA
961 CGAGGATGAA GAGGGTGAGG AGTTTGTGTT AGATTATGTG GAGCACCCCG GGCACGGTTG
1021 CAGGTCTTGT CATTATCACC GGAGGAATAC GGGGGACCCA GATATTATGT GTTCGCTTTG
1081 CTATATGAGG ACCTGTGGCA TGTTTGTCTA CAGTAAGTGA AAAATTATGG GCAGTGGGTG
1141 ATAGAGTGGT GGGTTTGGTG TGTAATTTT TTTTFTAATT TTTACAGTTT TGTGGTTTAA
1201 AGAATTTTGT ATTGTGATTT TTTAAAAGGT CCTGTGTCTG AACCTGAGCC TGAGCCCGAG
1261 CCAGAACCGG AGCCTGCAAG ACCTACCCGG CGTCCTAAAT TGGTGCCTGC TATCCTGAGA
1321 CGCCCGACAT CACCTGTGTC TAGAGAATGC AATAGTAGTA CGGATAGCTG TGACTCCGGT
1381 CCTTCTAACA CACCTCCTGA GATACACCCG GTGGTCCCGC TGTGCCCCAT TAAACCAGTT
1441 GCCGTGAGAG TTGGTGGGCG TCGCCAGGCT GTGGAATGTA TCGAGGACTT GCTTAACGAG
1501 TCTGGGCAAC CTTTGGACTT GAGCTGTAAA CGCCCAGGC CATAAGGTGT AAACCTGTGA
1561 TTGCGTGTGT GGTTAACGCC TTTGTTTGCT GAATGAGTTG ATGTAAGTTT AATAAAGGGT
1621 GAGATAATGT TTAACCTGCA TGGCGTGTTA AATGGGGCGG GGCTTAAAGG GTATATAATG
1681 CGCCGTGGGC TAATCTTGGT TACATCTGAC CTCATGGAGG CTTGGGAGTG TTTGGAAGAT
1741 TTTTCTGCTG TCGTAACTT GCTGGAACAG AGCTCTAACA GTACCTCTTG GTTTTGGAGG
1801 TTTCTGTGGG GCTCCTCCCA GGCAAAGTTA GTCTGCAGAA TTAAGGAGGA TTACAAGTGG
1861 GAATTTGAAG AGCTTTTGAA ATCCTGTGGT GAGCTGTTTG ATTCTTTGAA TCTGGGTCAC
1921 CAGGCGCTTT TCCAAGAGAA GGTCAATCAAG ACTTTGGATT TTTCCACACC GGGGCGCGCT
1981 GCGGCTGCTG TTGCTTTTTT GAGTTTTATA AAGGATAAAT GGAGCGAAGA AACCCTCTG
2041 AGCGGGGGGT ACCTGCTGGA TTTTCTGGCC ATGCATCTGT GGAGAGCGGT GGTGAGACAC
2101 AAGAATCGCC TGCTACTGTT GTCTTCCGTC CGCCCGCAA TAATACCGAC GGAGGAGCAA
2161 CAGCAGGAGG AAGCCAGGCG GCGGCGGCGG CAGGAGCAGA GCCCATGGAA CCCGAGAGCC
2221 GGCCTGGACC CTCGGGAATG AATGTTGTAC AGGTGGCTGA ACTGTTTCCA GAACTGAGAC
2281 GCATTTTAAC CATTACGAG GATGGGCAGG GGCTAAAGGG GGTAAAGAAG GAGCGGGGGG
2341 CTTCTGAGGC TACAGAGGAG GCTAGGAATC TAACTTTTAG CTTAATGACC AGACACCGTC
2401 CTGAGTGTGT TACTTTTCAG CAGATTAAGG ATAATTGCGC TAATGAGCTT GATCTGCTGG
2461 CGCAGAAGTA TTCCATAGAG CAGCTGACCA CTTACTGGCT GCAGCCAGGG GATGATTTTG
2521 AGGAGGCTAT TAGGGTATAT GCAAAGGTGG CACTTAGGCC AGATTGCAAG TACAAGATTA
2581 GCAAACCTTG AAATATCAGG AATTGTTGCT ACATTTCTGG GAACGGGGCC GAGGTGGAGA
2641 TAGATACGGA GGATAGGGTG GCCTTTAGAT GTAGCATGAT AAATATGTGG CCGGGGGTGC
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FIG.11A-1

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|      |            |            |             |            |            |            |
|------|------------|------------|-------------|------------|------------|------------|
| 2701 | TTGGCATGGA | CGGGGTGGTT | ATTATGAATG  | TGAGGTTTAC | TGGTCCCAAT | TTTAGCGGTA |
| 2761 | CGGTTTTCT  | GGCCAATACC | AATCTTATCC  | TACACGGTGT | AAGCTTCTAT | GGGTTTAACA |
| 2821 | ATACCTGTGT | GGAAGCCTGG | ACCGATGTAA  | GGGTTTCGGG | CTGTGCCTTT | TACTGCTGCT |
| 2881 | GGAAGGGGGT | GGTGTGTCGC | CCCAAAAGCA  | GGGCTTCAAT | TAAGAAATGC | CTGTTTGAAA |
| 2941 | GGTGTACCTT | GGGTATCCTG | TCTGAGGGTA  | ACTCCAGGGT | GCGCCACAAT | GTGGCCTCCG |
| 3001 | ACTGTGGTTG | CTTTATGCTA | GTGAAAAGCG  | TGGCTGTGAT | TAAGCATAAC | ATGGTGTGTG |
| 3061 | GCAACTGCGA | GGACAGGGCC | TCTCAGATGC  | TGACCTGCTC | GGACGGCAAC | TGTCACTTGC |
| 3121 | TGAAGACCAT | TCACGTAGCC | AGCCACTCTC  | GCAAGGCCTG | GCCAGTGTTT | GAGCACAACA |
| 3181 | TACTGACCCG | CTGTTCCCTG | CATTTGGGTA  | ACAGGAGGGG | GGTGTTCCTA | CCTTACCAAT |
| 3241 | GCAATTTGAG | TCACACTAAG | ATATTGCTTG  | AGCCCGAGAG | CATGTCCAAG | GTGAACCTGA |
| 3301 | ACGGGGTGTT | TGACATGACC | ATGAAGATCT  | GGAAGGTGCT | GAGGTACGAT | GAGACCCGCA |
| 3361 | CCAGGTGCAG | ACCCTGCGAG | TGTGGCGGTA  | AACATATTAG | GAACCAGCCT | GTGATGCTGG |
| 3421 | ATGTGACCGA | GGAGCTGAGG | CCCGATCACT  | TGGTGCTGGC | CTGCACCCGC | GCTGAGTTTG |
| 3481 | GCTCTAGCGA | TGAAGATACA | GATTGAGGTA  | CTGAAATGTG | TGGGCGTGCC | TTAAGGGTGG |
| 3541 | GAAAGAATAT | ATAAGGTGGG | GGTCTCATGT  | AGTTTTGTAT | CTGTTTTGCA | GCAGCCGCCG |
| 3601 | CCATGAGCGC | CAACTCGTTT | GATGGAAGCA  | TTGTGAGCTC | ATATTTGACA | ACGCGCATGC |
| 3661 | CCCCATGGGC | CGGGGTGCGT | CAGAATGTGA  | TGGGCTCCAG | CATTGATGGT | CGCCCCGTCC |
| 3721 | TGCCCGCAAA | CTCTACTACC | TTGACCTACG  | AGACCGTGTC | TGGAACGCCG | TTGGAGACTG |
| 3781 | CAGCCTCCGC | CGCCGCTTCA | GCCGCTGCAG  | CCACCGCCCG | CGGGATTGTG | ACTGACTTTG |
| 3841 | CTTTCCTGAG | CCCGCTTGCA | AGCAGTGCAG  | CTTCCCGTTC | ATCCGCCCGC | GATGACAAGT |
| 3901 | TGACGGCTCT | TTTGGCACAA | TTGGATTCTT  | TGACCCGGGA | ACTTAATGTC | GTTTCTCAGC |
| 3961 | AGCTGTTGGA | TCTGCGCCAG | CAGGTTTCTG  | CCCTGAAGGC | TTCCTCCCCT | CCCAATGCGG |
| 4021 | TTTAAACAT  | AAATAAAAC  | CAGACTCTGT  | TTGGATTTGG | ATCAAGCAAG | TGTCTTGCTG |
| 4081 | TCTTTATTTA | GGGGTTTTGC | GCGCGCGGTA  | GGCCCGGGAC | CAGCGGTCTC | GGTCGTTGAG |
| 4141 | GGTCCTGTGT | ATTTTTTCCA | GGACGTGGTA  | AAGGTGACTC | TGGATGTTCA | GATACATGGG |
| 4201 | CATAAGCCCG | TCTCTGGGGT | GGAGGTAGCA  | CCACTGCAGA | GCTTCATGCT | GCGGGGTGGT |
| 4261 | GTTGTAGATG | ATCCAGTCGT | AGCAGGAGCG  | CTGGGCGTGG | TGCCTAAAAA | TGTCTTTCAG |
| 4321 | TAGCAAGCTG | ATTGCCAGGG | GCAGGCCCTT  | GGTGTAAAGT | TTTACAAAGC | GGTTAAGCTG |
| 4381 | GGATGGGTGC | ATACGTGGGG | ATATGAGATG  | CATCTTGGAC | TGTATTTTTA | GGTTGGCTAT |
| 4441 | GTTCCAGCC  | ATATCCCTCC | GGGGATTTCAT | GTTGTGCAGA | ACCACCAGCA | CAGTGTATCC |
| 4501 | GGTGCACTTG | GGAAATTTGT | CATGTAGCTT  | AGAAGGAAAT | GCGTGGAAGA | ACTTGGAGAC |
| 4561 | GCCCTTGTGA | CCTCCAAGAT | TTTCCATGCA  | TTCGTCCATA | ATGATGGCAA | TGGGGCCACG |
| 4621 | GGCGGCGGCC | TGGGCGAAGA | TATTTCTGGG  | ATCACTAACG | TCATAGTTGT | GTTCCAGGAT |
| 4681 | GAGATCGTCA | TAGGCCATTT | TTACAAAGCG  | CGGGCGGAGG | GTGCCAGACT | GCGGTATAAT |
| 4741 | GGTTCCATCC | GGCCAGGGG  | CGTAGTTACC  | CTCACAGATT | TGCATTTCCC | ACGCTTTGAG |
| 4801 | TTCAGATGGG | GGGATCATGT | CTACCTGCGG  | GGCGATGAAG | AAAACCGTTT | CCGGGGTAGG |
| 4861 | GGAGATCAGC | TGGGAAGAAA | GCAGGTTCTT  | AAGCAGCTGC | GACTTACCGC | AGCCGGTGGG |
| 4921 | CCCGTAAATC | ACACCTATTA | CCGGCTGCAA  | CTGGTAGTTA | AGAGAGCTGC | AGCTGCCGTC |
| 4981 | ATCCCTGAGC | AGGGGGGCCA | CTTCGTTAAG  | CATGTCCCTG | ACTTGCATGT | TTTCCCTGAC |
| 5041 | CAAATCCGCC | AGAAGGCGCT | CGCCGCCCAG  | CGATAGCAGT | TCTTGCAAGG | AAGCAAAGTT |
| 5101 | TTTCAACGGT | TTGAGGCCGT | CCGCCGTAGG  | CATGCTTTTG | AGCGTTTGAC | CAAGCAGTTC |
| 5161 | CAGGCGGTCC | CACAGCTCGG | TCACGTGCTC  | TACGGCATCT | CGATCCAGCA | TATCTCCTCG |
| 5221 | TTTCGCGGGT | TGGGGCGGCT | TTCGCTGTAC  | GGCAGTAGTC | GGTGCTCGTC | CAGACGGGCC |
| 5281 | AGGGTCATGT | CTTTCACGG  | GCGCAGGGTC  | CTCGTCAGCG | TAGTCTGGGT | CACGGTGAAG |
| 5341 | GGGTGCGCTC | CGGGTTGCGC | GCTGGCCAGG  | GTGCGCTTGA | GGCTGGTCCT | GCTGGTGCTG |

FIG. 11A-2



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5401 AAGCGCTGCC GGTCTTCGCC CTGCGCGTCG GCCAGGTAGC ATTTGACCAT GGTGTCATAG  
 5461 TCCAGCCCCT CCGCGGCGTG GCCCTTGGCG CGCAGCTTGC CTTTGGAGGA GGCGCCGCAC  
 5521 GAGGGGCAGT GCAGACTTTT AAGGGCGTAG AGCTTGGGCG CGAGAAATAC CGATTCCGGG  
 5581 GAGTAGGCAT CCGCGCCGCA GGCCCCGCAG ACGGTCTCGC ATTCCACGAG CCAGGTGAGC  
 5641 TCTGGCCGTT CGGGGTCAAA AACCAGGTTT CCCCCATGCT TTTTGATGCG TTTCTTACCT  
 5701 CTGGTTTCCA TGAGCCGGTG TCCACGCTCG GTGACGAAAA GGCTGTCCGT GTCCCCGTAT  
 5761 ACAGACTTGA GAGGCCTGTC CTCGAGCGGT GTTCCGCGGT CCTCCTCGTA TAGAAACTCG  
 5821 GACCACTCTG AGACGAAGGC TCGCGTCCAG GCCAGCACGA AGGAGGCTAA GTGGGAGGGG  
 5881 TAGCGGTCGT TGTCCACTAG GGGGTCCACT CGCTCCAGGG TGTGAAGACA CATGTCGCCC  
 5941 TCTTCGGCAT CAAGGAAGGT GATTGGTTTA TAGGTGTAGG CCACGTGACC GGGTGTTTCT  
 6001 GAAGGGGGGC TATAAAAGGG GGTGGGGGCG CGTTCGTCCT CACTCTCTTC CGCATCGCTG  
 6061 TCTGCGAGGG CCAGCTGTTG GGGTGAGTAC TCCCTCTCAA AAGCGGGCAT GACTTCTGCG  
 6121 CTAAGATTGT CAGTTTCCAA AAACGAGGAG GATTTGATAT TCACCTGGCC CGCGGTGATG  
 6181 CCTTTGAGGG TGGCCGCGTC CATCTGGTCA GAAAAGACAA TCTTTTGTG GTCAAGCTTG  
 6241 GTGGCAAACG ACCCGTAGAG GCGCTTGGAC AGCAACTTGG CGATGGAGCG CAGGGTTTGG  
 6301 TTTTGTGCG GATCGGCGCG CTCCTTGGCC GCGATGTTTA GCTGCACGTA TTCGCGCGCA  
 6361 ACGCACCGCC ATTCGGGAAA GACGGTGGTG CGCTCGTCGG GCACTAGGTG CACGCGCCAA  
 6421 CCGCGGTTGT GCAGGGTGAC AAGGTCAACG CTGGTGGCTA CCTCTCCGCG TAGGCGCTCG  
 6481 TTGGTCCAGC AGAGGCGGCC GCCCTTGC GC GAGCAGAATG GCGGTAGTGG GTCTAGCTGC  
 6541 GTCTCGTCCG GGGGGTCTGC GTCCACGGTA AAGACCCCGG GCAGCAGGCG CGCGTCGAAG  
 6601 TAGTCTATCT TGCATCCTTG CAAGTCTAGC GCCTGCTGCC ATGCGCGGGC GGCAAGCGCG  
 6661 CGCTCGTATG GGTTGAGTGG GGGACCCCAT GGCATGGGGT GGGTGAGCGC GGAGGCGTAC  
 6721 ATGCCGAAA TGTCGTAAAC GTAGAGGGGC TCTCTGAGTA TTCCAAGATA TGTAGGGTAG  
 6781 CATCTTCCAC CGCGGATGCT GGCGCGCACG TAATCGTATA GTTCGTGCGA GGGAGCGAGG  
 6841 AGGTCGGGAC CGAGGTTGCT ACGGGCGGGC TGCTCTGCTC GGAAGACTAT CTGCCTGAAG  
 6901 ATGGCATGTG AGTTGGATGA TATGGTTGGA CGCTGGAAGA CGTTGAAGCT GGCCTCTGTG  
 6961 AGACCTACCG CGTCACGCAC GAAGGAGGCG TAGGAGTCGC GCAGCTTGTT GACCAGCTCG  
 7021 GCGGTGACCT GCACGTCTAG GGCGCAGTAG TCCAGGGTTT CCTTGATGAT GTCATACTTA  
 7081 TCCTGTCCCT TTTTTTCCA CAGCTCGCGG TTGAGGACAA ACTCTTCGCG GTCTTTCCAG  
 7141 TACTCTTGA TCGGAAACCC GTCGGCCTCC GAACGGTAAG AGCCTAGCAT GTAGAACTGG  
 7201 TTGACGGCCT GGTAGGCGCA GCATCCCTTT TCTACGGGTA GCGCGTATGC CTGCGCGGCC  
 7261 TTCCGGAGCG AGGTGTGGGT GAGCGCAAAG GTGTCCCTAA CCATGACTTT GAGGTACTGG  
 7321 TATTTGAAGT CAGTGTGCTC GCATCCGCCC TGCTCCCAGA GCAAAAAGTC CGTGCGCTTT  
 7381 TTGGAACGCG GGTTTGGCAG GGCGAAGGTG ACATCGTTGA AGAGTATCTT TCCCGCGCGA  
 7441 GGCATAAAGT TGC GTGTGAT GCGGAAGGGT CCCGGCACCT CGGAACGGTT GTTAATTACC  
 7501 TGGGCGGCGA GCACGATCTC GTCAAAGCCG TTGATGTTGT GGCCACAAT GTAAAGTTCC  
 7561 AAGAAGCGCG GGATGCCCTT GATGGAAGGC AATTTTTTAA GTTCCTCGTA GGTGAGCTCT  
 7621 TCAGGGGAGC TGAGCCCGTG CTCTGAAAGG GCCAGTCTG CAAGATGAGG GTTGGAAGCG  
 7681 ACGAATGAGC TCCACAGGTC ACGGGCCATT AGCATTTGCA GGTGGTTCGCG AAAGGTCCTA  
 7741 AACTGGCGAC CTATGGCCAT TTTTCTGGG GTGATGCAGT AGAAGGTAAG CGGGTCTTGT  
 7801 TCCCAGCGGT CCCATCCAAG GTCCGCGGCT AGGTCTCGCG CGGCGGTAC TAGAGGCTCA  
 7861 TCTCCGCGGA ACTTCATGAC CAGCATGAAG GGCACGAGCT GCTTCCCAA GGCCCCATC  
 7921 CAAGTATAGG TCTCTACATC GTAGGTGACA AAGAGACGCT CGGTGCGAGG ATGCGAGCCG  
 7981 ATCGGGAAGA ACTGGATCTC CCGCCACCAG TTGGAGGAGT GGCTGTTGAT GTGGTGAAAG  
 8041 TAGAAGTCCC TGCGACGGGC CGAACACTCG TGCTGGCTTT TGTA AAAACG TGCGCAGTAC

FIG. 11A-3

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|       |            |             |            |             |            |            |
|-------|------------|-------------|------------|-------------|------------|------------|
| 8101  | TGGCAGCGGT | GCACGGGCTG  | TACATCCTGC | ACGAGGTTGA  | CCTGACGACC | GCGCACAAGG |
| 8161  | AAGCAGAGTG | GGAATTTGAG  | CCCCTCGCCT | GGCGGGTTTG  | GCTGGTGGTC | TTCTACTTCG |
| 8221  | GCTGCTTGTC | CTTGACCGTC  | TGGCTGCTCG | AGGGGAGTTA  | CGGTGGATCG | GACCACCACG |
| 8281  | CCGCGCGAGC | CCAAAGTCCA  | GATGTCCGCG | CGCGGCGGTC  | GGAGCTTGAT | GACAACATCG |
| 8341  | CGCAGATGGG | AGCTGTCCAT  | GGTCTGGAGC | TCCC GCGGCG | TCAGGTCAGG | CGGGAGCTCC |
| 8401  | TGCAGGTTTA | CCTCGCATAG  | CCGGGTCAGG | GCGCGGGCTA  | GGTCCAGGTG | ATACCTGATT |
| 8461  | TCCAGGGGCT | GGTTGGTGGC  | GGCGTCGATG | GCTTGCAAGA  | GGCCGCATCC | CCGCGGCGCG |
| 8521  | ACTACGGTAC | CGCGCGGCGG  | GCGGTGGGCC | GCGGGGGTGT  | CCTTGGATGA | TGCATCTAAA |
| 8581  | AGCGGTGACG | CGGGCGGGCC  | CCCGGAGGTA | GGGGGGGCTC  | GGGACCCGCC | GGGAGAGGGG |
| 8641  | GCAGGGGCAC | GTCGGCGCCG  | CGCGCGGGCA | GGAGCTGGTG  | CTGCGCGCGG | AGGTTGCTGG |
| 8701  | CGAACGCGAC | GACGCGGCGG  | TTGATCTCCT | GAATCTGGCG  | CCTCTGCGTG | AAGACGACGG |
| 8761  | GCCCGGTGAG | CTTGAACCTG  | AAAGAGAGTT | CGACAGAATC  | AATTTGCGTG | TCGTTGACGG |
| 8821  | CGGCCTGGCG | CAAAATCTCC  | TGCACGTCTC | CTGAGTTGTC  | TTGATAGGCG | ATCTCGGCCA |
| 8881  | TGAACTGCTC | GATCTCTTCC  | TCCTGGAGAT | CTCCGCGTCC  | GGCTCGCTCC | ACGGTGGCGG |
| 8941  | CGAGGTCGTT | GGAGATGCGG  | GCCATGAGCT | GCGAGAAGGC  | GTTGAGGCCT | CCCTCGTTCC |
| 9001  | AGACGCGGCT | GTAGACCACG  | CCCCCTTCGG | CATCGCGGGC  | GCGCATGACC | ACCTGCGCGA |
| 9061  | GATTGAGCTC | CACGTGCCGG  | GCGAAGACGG | CGTAGTTTCG  | CAGGCGCTGA | AAGAGGTAGT |
| 9121  | TGAGGGTGGT | GGCGGTGTGT  | TCTGCCACGA | AGAAGTACAT  | AACCCAGCGC | CGCAACGTGG |
| 9181  | ATTCGTTGAT | ATCCCCCAAG  | GCCTCAAGGC | GCTCCATGGC  | CTCGTAGAAG | TCCACGGCGA |
| 9241  | AGTTGAAAAA | CTGGGAGTTG  | CGCGCCGACA | CGGTTAACTC  | CTCCTCCAGA | AGACGGATGA |
| 9301  | GCTCGGCGAC | AGTGTCGCGC  | ACCTCGCGCT | CAAAGGCTAC  | AGGGGCCTCT | TCTTCTTCTT |
| 9361  | CAATCTCCTC | TTCCATAAGG  | GCCTCCCCTT | CTTCTTCTTC  | TGGCGGCGGT | GGGGGAGGGG |
| 9421  | GGACACGGCG | GCGACGACGG  | CGCACCGGGA | GGCGGTCGAC  | AAAGCGCTCG | ATCATCTCCC |
| 9481  | CGCGGCGACG | GCGCATGGTC  | TCGGTGACGG | CGCGGCCGTT  | CTCGCGGGGG | GCGAGTTGGA |
| 9541  | AGACGCCGCC | CGTCATGTCC  | CGGTTATGGG | TTGGCGGGGG  | GCTGCCGTGC | GGCAGGGATA |
| 9601  | CGGCGCTAAC | GATGCATCTC  | AACAATTGTT | GTGTAGGTAC  | TCCGCCACCG | AGGGACCTGA |
| 9661  | GCGAGTCCGC | ATCGACCGGA  | TCGGAAAACC | TCTCGAGAAA  | GGCGTCTAAC | CAGTCACAGT |
| 9721  | CGCAAGGTAG | GCTGAGCACC  | GTGGCGGGCG | GCAGCGGGCG  | GCGGTCGGGG | TTGTTTCTGG |
| 9781  | CGGAGGTGCT | GCTGATGATG  | TAATTAAGT  | AGGCGGTCTT  | GAGACGGCGG | ATGGTCGACA |
| 9841  | GAAGCACCAT | GTCCTTGGGT  | CCGGCCTGCT | GAATGCGCAG  | GCGGTCGGCC | ATGCCCCAGG |
| 9901  | CTTCGTTTTG | ACATCGGCGC  | AGGTCTTTGT | AGTAGTCTTG  | CATGAGCCTT | TCTACCGGCA |
| 9961  | CTTCTTCTTC | TCCTTCCTCT  | TGTCCTGCAT | CTCTTGTCATC | TATCGCTGCG | GCGGCGGCGG |
| 10021 | AGTTTGGCCG | TAGGTGGCGC  | CCTCTTCCTC | CCATGCGTGT  | GACCCCGAAG | CCCCTCATCG |
| 10081 | GCTGAAGCAG | GGCCAGGTCTG | GCGACAACGC | GCTCGGCTAA  | TATGGCCTGC | TGCACCTGCG |
| 10141 | TGAGGGTAGA | CTGGAAGTCG  | TCCATGTCCA | CAAAGCGGTG  | GTATGCGCCC | GTGTTGATGG |
| 10201 | TGTAAGTGCA | GTTGGCCATA  | ACGGACCAGT | TAACGGTCTG  | GTGACCCGGC | TGCGAGAGCT |
| 10261 | CGGTGTACCT | GAGACGCGAG  | TAAGCCCTTG | AGTCAAAGAC  | GTAGTCGTTG | CAAGTCCGCA |
| 10321 | CCAGGTACTG | GTATCCACC   | AAAAAGTGCG | GCGGCGGCTG  | GCGGTAGAGG | GGCCAGCGTA |
| 10381 | GGGTGGCCGG | GGCTCCGGGG  | GCGAGGTCTT | CCAACATAAG  | GCGATGATAT | CCGTAGATGT |
| 10441 | ACCTGGACAT | CCAGGTGATG  | CCGGCGGCGG | TGGTGGAGGC  | GCGCGGAAAG | TCACGGACGC |
| 10501 | GGTTCCAGAT | GTTGCGCAGC  | GGCAAAAAGT | GCTCCATGGT  | CGGGACGCTC | TGGCCGGTCA |
| 10561 | GGCGCGCGCA | GTCGTTGACG  | CTCTAGACCG | TGCAAAAGGA  | GAGCCTGTAA | GCGGGCACTC |
| 10621 | TTCCGTGGTC | TGGTGGATAA  | ATTCGCAAGG | GTATCATGGC  | GGACGACCGG | GGTTCGAACC |
| 10681 | CCGGATCCGG | CCGTCCGCCG  | TGATCCATGC | GGTTACCGCC  | CGCGTGTCGA | ACCCAGGTGT |
| 10741 | GCGACGTCAG | ACAACGGGGG  | AGCGCTCCTT | TTGGCTTCCT  | TCCAGGCGCG | GCGGATGCTG |

FIG.11A-4

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|       |             |            |            |             |            |             |
|-------|-------------|------------|------------|-------------|------------|-------------|
| 10801 | CGCTAGCTTT  | TTTGGCCACT | GGCCGCGCGC | GGCGTAAGCG  | GTTAGGCTGG | AAAGCGAAAG  |
| 10861 | CATTAAGTGG  | CTCGCTCCCT | GTAGCCGGAG | GGTTATTTTC  | CAAGGGTTGA | GTCGCGGGAC  |
| 10921 | CCCCGGTTTCG | AGTCTCGGGC | CGGCCGGACT | GCGGCGAACG  | GGGGTTTGCC | TCCCCGTCAT  |
| 10981 | GCAAGACCCC  | GCTTGCAAAT | TCCTCCGGAA | ACAGGGACGA  | GCCCCTTTTT | TGCTTTTTCCC |
| 11041 | AGATGCATCC  | GGTGCTGCGG | CAGATGCGCC | CCCCTCCTCA  | GCAGCGGCAA | GAGCAAGAGC  |
| 11101 | AGCGGCAGAC  | ATGCAGGGCA | CCCTCCCCTT | CTCCTACCGC  | GTCAGGAGGG | GCAACATCCG  |
| 11161 | CGGCTGACGC  | GGCGGCAGAT | GGTGATTACG | AACCCCCGCG  | GCGCCGGACC | CGGCACTACT  |
| 11221 | TGGACTTGGA  | GGAGGGCGAG | GGCCTGGCGC | GGCTAGGAGC  | GCCCTCTCCT | GAGCGACACC  |
| 11281 | CAAGGGTGCA  | GCTGAAGCGT | GACACGCGCG | AGGCGTACGT  | GCCGCGGCAG | AACCTGTTTC  |
| 11341 | GCGACCGCGA  | GGGAGAGGAG | CCCGAGGAGA | TGCGGGATCG  | AAAGTTCCAT | GCAGGGCGCG  |
| 11401 | AGTTGCGGCA  | TGGCCTGAAC | CGCGAGCGGT | TGCTGCGCGA  | GGAGGACTTT | GAGCCCCGACG |
| 11461 | CGCGGACCGG  | GATTAGTCCC | GCGCGCGCAC | ACGTGGCGGC  | CGCCGACCTG | GTAACCGCGT  |
| 11521 | ACGAGCAGAC  | GGTGAACCAG | GAGATTAAC  | TTCAAAAAAG  | CTTTAACAAC | CACGTGCGCA  |
| 11581 | CGCTTGTTGGC | GCGCGAGGAG | GTGGCTATAG | GACTGATGCA  | TCTGTGGGAC | TTTGTAAAGCG |
| 11641 | CGCTGGAGCA  | AAACCCAAAT | AGCAAGCCGC | TCATGGCGCA  | GCTGTTCCCT | ATAGTGCAGC  |
| 11701 | ACAGCAGGGA  | CAACGAGGCA | TTCAGGGATG | CGCTGCTAAA  | CATAGTAGAG | CCCGAGGGCC  |
| 11761 | GCTGGCTGCT  | CGATTTGATA | AACATTCTGC | AGAGCATAGT  | GGTGCAGGAG | CGCAGCTTGA  |
| 11821 | GCCTGGCTGA  | CAAGGTGGCC | GCCATTAAC  | ATTCCATGCT  | CAGTCTGGGC | AAGTTTTACG  |
| 11881 | CCCGCAAGAT  | ATACCATACC | CCTTACGTTT | CCATAGACAA  | GGAGGTAAAG | ATCGAGGGGT  |
| 11941 | TCTACATGCG  | CATGGCGCTG | AAGGTGCTTA | CCTTGAGCGA  | CGACCTGGGC | GTTTATCGCA  |
| 12001 | ACGAGCGCAT  | CCACAAGGCC | GTGAGCGTGA | GCCGGCGGCG  | CGAGCTCAGC | GACCGCGAGC  |
| 12061 | TGATGCACAG  | CCTGCAAAGG | GCCCTGGCTG | GCACGGGCAG  | CGGCGATAGA | GAGGCCGAGT  |
| 12121 | CCTACTTTGA  | CGCGGGCGCT | GACCTGCGCT | GGGCCCCAAG  | CCGACGCGCC | CTGGAGGCAG  |
| 12181 | CTGGGGCCGG  | ACCTGGGCTG | GCGGTGGCAC | CCGCGCGCGC  | TGGCAACGTC | GGCGGCGTGG  |
| 12241 | AGGAATATGA  | CGAGGACGAT | GAGTACGAGC | CAGAGGACGG  | CGAGTACTAA | GCGGTGATGT  |
| 12301 | TTCTGATCAG  | ATGATGCAAG | ACGCAACGGA | CCCGGCGGTG  | CGGGCGGCGC | TGCAGAGCCA  |
| 12361 | GCCGTCCGGC  | CTTAACCTCA | CGGACGACTG | GCGCCAGGTC  | ATGGACCGCA | TCATGTCGCT  |
| 12421 | GACTGCGCGC  | AACCCTGACG | CGTTCCGGCA | GCAGCCGCAG  | GCCAACCGGC | TCTCCGCAAT  |
| 12481 | TCTGGAAGCG  | GTGGTCCCGG | CGCGCGCAAA | CCCCACGCAC  | GAGAAGGTGC | TGGCGATCGT  |
| 12541 | AAACGCGCTG  | GCCGAAAACA | GGGCCATCCG | GCCCCGATGAG | GCCGGCCTGG | TCTACGACGC  |
| 12601 | GCTGCTTCAG  | CGCGTGGCTC | GTTACAACAG | CAGCAACGTG  | CAGACCAACC | TGGACCGGCT  |
| 12661 | GGTGGGGGAT  | GTGCGCGAGG | CCGTGGCGCA | GCGTGAGCGC  | GCGCAGCAGC | AGGGCAACCT  |
| 12721 | GGGCTCCATG  | GTTGCACTAA | ACGCCTTCCT | GAGTACACAG  | CCCGCCAACG | TGCCGCGGGG  |
| 12781 | ACAGGAGGAC  | TACACCAACT | TTGTGAGCGC | ACTGCGGCTA  | ATGGTGACTG | AGACACCGCA  |
| 12841 | AAGTGAGGTG  | TATCAGTCCG | GGCCAGACTA | TTTTTTCCAG  | ACCAGTAGAC | AAGGCCTGCA  |
| 12901 | GACCGTAAAC  | CTGAGCCAGG | CTTTCAAGAA | CTTGCAAGGG  | CTGTGGGGGG | TGCGGGCTCC  |
| 12961 | CACAGGCGAC  | CGCGCGACCG | TGTCTAGCTT | GCTGACGCCC  | AACTCGCGCC | TGTTGCTGCT  |
| 13021 | GCTAATAGCG  | CCCTTCACGG | ACAGTGGCAG | CGTGTCCCGG  | GACACATACC | TAGGTCACTT  |
| 13081 | GCTGACACTG  | TACCGCGAGG | CCATAGGTCA | GGCGCATGTG  | GACGAGCATA | CTTTCCAGGA  |
| 13141 | GATTACAAGT  | GTTAGCCGCG | CGCTGGGGCA | GGAGGACACG  | GGCAGCCTGG | AGGCAACCCT  |
| 13201 | GAACCTACCTG | CTGACCAACC | GGCGGCAAAA | AATCCCCTCG  | TTGCACAGTT | TAAACAGCGA  |
| 13261 | GGAGGAGCGC  | ATTTTGCGCT | ATGTGCAGCA | GAGCGTGAGC  | CTTAACCTGA | TGCGCGACGG  |
| 13321 | GGTAACGCC   | AGCGTGGCGC | TGGACATGAC | CGCGCGCAAC  | ATGGAACCGG | GCATGTATGC  |
| 13381 | CTCAAACCGG  | CCGTTTATCA | ATCGCCTAAT | GGACTACTTG  | CATCGCGCGG | CCGCCGTGAA  |
| 13441 | CCCCGAGTAT  | TTCACCAATG | CCATCTTGAA | CCCGCACTGG  | CTACCGCCCC | CTGGTTTCTA  |

FIG. 11A-5

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|       |            |            |             |             |            |            |
|-------|------------|------------|-------------|-------------|------------|------------|
| 13501 | CACCGGGGGA | TTCGAGGTGC | CCGAGGGTAA  | CGATGGATTG  | CTCTGGGACG | ACATAGACGA |
| 13561 | CAGCGTGTTT | TCCCCGCAAC | CGCAGACCCCT | GCTAGAGTTG  | CAACAACGCG | AGCAGGCAGA |
| 13621 | GGCGGCGCTG | CGAAAGGAAA | GCTTCCGCAG  | GCCAAGCAGC  | TTGTCCGATC | TAGGCGCTGC |
| 13681 | GGCCCCGCGG | TCAGATGCTA | GTAGCCCAT   | TCCAAGCTTG  | ATAGGGTCTC | TTACCAGCAC |
| 13741 | TCGCACCACC | CGCCCGCGCC | TGCTGGGCGA  | GGAGGAGTAC  | CTAAACAAC  | CGCTGCTGCA |
| 13801 | GCCGCAGCGC | GAAAAGAACC | TGCCTCCGGC  | GTTTCCCAAC  | AACGGGATAG | AGAGCCTAGT |
| 13861 | GGACAAGATG | AGTAGATGGA | AGACGTATGC  | GCAGGAGCAC  | AGGGATGTGC | CCGGCCCGCG |
| 13921 | CCCGCCCAAC | CGTCGTCAAA | GGCACGACCG  | TCAGCGGGGT  | CTGGTGTGGG | AGGACGATGA |
| 13981 | CTCGGCAGAC | GACAGCAGCG | TCTTGGATTT  | GGGAGGGAGT  | GGCAACCCGT | TTGCACACCT |
| 14041 | TCGCCCCAGG | CTGGGGAGAA | TGTTTTAAAA  | AAAGCATGAT  | GCAAAATAAA | AAACTCACCA |
| 14101 | AGGCCATGGC | ACCGAGCGTT | GGTTTTCTTG  | TATTCCTT    | AGTATGCGGC | GCGCGGCGAT |
| 14161 | GTATGAGGAA | GGTCCTCCTC | CCTCCTACGA  | GAGCGTGGTG  | AGCGCGGCGC | CAGTGGCGGC |
| 14221 | GGCGCTGGGT | TCACCCTTCG | ATGCTCCCCT  | GGACCCGCCG  | TTCGTGCCTC | CGCGGTACCT |
| 14281 | GCGGCCTACC | GGGGGGAGAA | ACAGCATCCG  | TTACTCTGAG  | TTGGCACCCC | TATTCGACAC |
| 14341 | CACCCGTGTG | TACCTTGTGG | ACAACAAGTC  | AACGGATGTG  | GCATCCCTGA | ACTACCAGAA |
| 14401 | CGACCACAGC | AACTTTCTAA | CCACGGTCAT  | TCAAAACAAT  | GACTACAGCC | CGGGGGAGGC |
| 14461 | AAGCACACAG | ACCATCAATC | TTGACGACCG  | GTCGCACTGG  | GGCGGCGACC | TGAAAACCAT |
| 14521 | CCTGCATACC | AACATGCCAA | ATGTGAACGA  | GTTTCATGTT  | ACCAATAAGT | TTAAGGCGCG |
| 14581 | GGTGATGGTG | TCGCGCTCGC | TTACTAAGGA  | CAAACAGGTG  | GAGCTGAAAT | ACGAGTGGGT |
| 14641 | GGAGTTCACG | CTGCCCAGAG | GCAACTACTC  | CGAGACCATG  | ACCATAGACC | TTATGAACAA |
| 14701 | CGCGATCGTG | GAGCACTACT | TGAAAGTGGG  | CAGGCAGAAC  | GGGGTTCTGG | AAAGCGACAT |
| 14761 | CGGGGTAAAG | TTTGACACCC | GCAACTTCAG  | ACTGGGGTTT  | GACCCAGTCA | CTGGTCTTGT |
| 14821 | CATGCCTGGG | GTATATACAA | ACGAAGCCTT  | CCATCCAGAC  | ATCATTTCG  | TGCCAGGATG |
| 14881 | CGGGGTGGAC | TTACCCACA  | GCCGCCTGAG  | CAACTTGTTG  | GGCATCCGCA | AGCGGCAACC |
| 14941 | CTTCCAGGAG | GGCTTTAGGA | TCACCTACGA  | TGACCTGGAG  | GGTGGTAACA | TTCCCGCACT |
| 15001 | GTTGGATGTG | GACGCCTACC | AGGCAAGCTT  | GAAAGATGAC  | ACCGAACAGG | GCGGGGGTGG |
| 15061 | CGCAGGCGGC | GGCAACAACA | GTGGCAGCGG  | CGCGGAAGAG  | AACTCCAACG | CGGCAGCTGC |
| 15121 | GGCAATGCAG | CCGGTGGAGG | ACATGAACGA  | TCATGCCATT  | CGCGGCGACA | CCTTTGCCAC |
| 15181 | ACGGGCGGAG | GAGAAGCGCG | CTGAGGCCGA  | GGCAGCGGCC  | GAAGCTGCCG | CCCCCGCTGC |
| 15241 | GGAGGCTGCA | CAACCCGAGG | TCGAGAAGCC  | TCAGAAGAAA  | CCGGTGATTA | AACCCCTGAC |
| 15301 | AGAGGACAGC | AAGAAACGCA | GTTACAACCT  | AATAAGCAAT  | GACAGCACCT | TCACCCAGTA |
| 15361 | CCGCAGCTGG | TACCTTGCAT | ACAACCTACG  | CGACCCCTCAG | GCCGGGATCC | GCTCATGGAC |
| 15421 | CCTGCTTTGC | ACTCCTGACG | TAACCTGCGG  | CTCGGAGCAG  | GTATACTGGT | CGTTGCCCGA |
| 15481 | CATGATGCAA | GACCCCGTGA | CCTTCCGCTC  | CACGCGCCAG  | ATCAGCAACT | TTCCGGTGGT |
| 15541 | GGGCGCCGAG | CTGTTGCCCG | TGCACTCCAA  | GAGCTTCTAC  | AACGACCAGG | CCGTCTACTC |
| 15601 | CCAGCTCATC | CGCCAGTTTA | CCTCTCTGAC  | CCACGTGTTT  | AATCGCTTTC | CCGAGAACCA |
| 15661 | GATTTTGGCG | CGCCCGCCAG | CCCCCACCAT  | CACCACCGTC  | AGTGAAAACG | TTCTGTCTCT |
| 15721 | CACAGATCAC | GGGACGCTAC | CGCTGCGCAA  | CAGCATCGGA  | GGAGTCCAGC | GAGTGACCAT |
| 15781 | TACTGACGCC | AGACGCCGCA | CCTGCCCTTA  | CGTTTACAAG  | GCCCTGGGCA | TAGTCTCGCC |
| 15841 | GCGGTCCTA  | TCGAGCCGCA | CTTTTGTAGC  | AAGCATGTCC  | ATCCTTATAT | CGCCAGCAA  |
| 15901 | TAACACAGGC | TGGGGCCTGC | GCTTCCCAAG  | CAAGATGTTT  | GGCGGGGCCA | AGAAGCGCTC |
| 15961 | CGACCAACAC | CCAGTGCGCG | TGCGCGGGCA  | CTACCGCGCG  | CCCTGGGGCG | CGCACAAACG |
| 16021 | CGGCCGCACT | GGGCGCACCA | CCGTGCATGA  | CGCCATCGAC  | GCGGTGGTGG | AGGAGGCGCG |
| 16081 | CAACTACACG | CCCACGCCGC | CGCCAGTGTC  | CACCGTGGAC  | GCGGCCATTG | AGACCGTGGT |
| 16141 | GCGCGGAGCC | CGGCGCTACG | CTAAAATGAA  | GAGACGGCGG  | AGGCGCGTAG | CACGTCGCCA |

FIG. 11A-6

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|       |             |            |             |             |            |            |
|-------|-------------|------------|-------------|-------------|------------|------------|
| 16201 | CCGCCGCCGA  | CCCGGCACTG | CCGCCCAACG  | CGCGGCGGCG  | GCCCTGCTTA | ACCGCGCACG |
| 16261 | TCGCACCGGC  | CGACGGGCGG | CCATGCGAGC  | CGCTCGAAGG  | CTGGCCGCGG | GTATTGTCAC |
| 16321 | TGTGCCCCCC  | AGGTCCAGGC | GACGAGCGGC  | CGCCGCAGCA  | GCCGCGGCCA | TTAGTGCTAT |
| 16381 | GA CTCAGGGT | CGCAGGGGCA | ACGTGTACTG  | GGTGCGCGAC  | TCGGTTAGCG | GCCTGCGCGT |
| 16441 | GCCCGTGCGC  | ACCCGCCCCC | CGCGCAACTA  | GATTGCAATA  | AAAACTACT  | TAGACTCGTA |
| 16501 | CTGTTGTATG  | TATCCAGCGG | CGGCGGCGCG  | CATCGAAGCT  | ATGTCCAAGC | GCAAAATCAA |
| 16561 | AGAAGAGATG  | CTCCAGGTCA | TCGCGCCGGA  | GATCTATGGC  | CCCCGAAGA  | AGGAAGAGCA |
| 16621 | GGATTACAAG  | CCCCGAAAGC | TAAAGCGGGT  | CAAAAAGAAA  | AAGAAAGATG | ATGATGATGA |
| 16681 | TGAAC TTGAC | GACGAGGTGG | AACTGTTGCA  | CGCGACCGCG  | CCCAGGCGAC | GGGTACAGTG |
| 16741 | GAAAGGT CGA | CGCGTAAGAC | GTGTTTTGCG  | ACCCGGCACC  | ACCGTAGTCT | TTACGCCCGG |
| 16801 | TGAGCGCTCC  | ACCCGCACCT | ACAAGCGCGT  | GTATGATGAG  | GTGTACGGCG | ACGAGGACCT |
| 16861 | GCTTGAGCAG  | GCCAACGAGC | GCCTCGGGGA  | GTTTGCCCTAC | GGAAAGCGGC | ATAAGGACAT |
| 16921 | GCTGGCGTTG  | CCGCTGGACG | AGGGCAACCC  | AACACCTAGC  | CTAAAGCCCG | TGACACTGCA |
| 16981 | GCAGGTGCTG  | CCCGCGCTTG | CACCGTCCGA  | AGAAAAGCGC  | GGCCTAAAGC | GCGAGTCTGG |
| 17041 | TGACTTG GCA | CCCACCGTGC | AGCTGATGGT  | ACCCAAGCGT  | CAGCGACTGG | AAGATGTCTT |
| 17101 | GGAAAAHATG  | ACCGTGGAGC | CTGGGCTGGA  | GCCCGAGGTC  | CGCGTGCGGC | CAATCAAGCA |
| 17161 | GGTGGCACCG  | GGACTGGGCG | TGCAGACCGT  | GGACGTT CAG | ATACCCACCA | CCAGTAGCAC |
| 17221 | TAGTATTGCC  | ACTGCCACAG | AGGGCATGGA  | GACACAAACG  | TCCCCGGTTG | CCTCGGCGGT |
| 17281 | GGCAGATGCC  | GCGGTGCAGG | CGGCCGCTGC  | GGCCGCGTCC  | AAGACCTCTA | CGGAGGTGCA |
| 17341 | AACGGACCCG  | TGGATGTTTC | GTGTTTCAGC  | CCCCCGGCGT  | CCGCGCCGTT | CAAGGAAGTA |
| 17401 | CGGCGCCGCC  | AGCGCGCTAC | TGCCCGAATA  | TGCCCTACAT  | CCTTCCATCG | CGCCTACCCC |
| 17461 | CGGCTATCGT  | GGCTACACCT | ACCGCCCCAG  | AAGACGAGCA  | ACTACCCGAC | GCCGAACCAC |
| 17521 | CACTGGAACC  | CGCCGCCGCC | GTCGCCGTCG  | CCAGCCCGTG  | CTGGCCCCGA | TTTCCGTGCG |
| 17581 | CAGGGTGGCT  | CGCGAAGGAG | GCAGGACCC T | GGTGCTGCCA  | ACAGCGCGCT | ACCACCCCAG |
| 17641 | CATCGTTTAA  | AAGCCGGTCT | TTGTGGTTCT  | TGCAGATATG  | GCCCTCACCT | GCCGCCCTCG |
| 17701 | TTTCCCGGTG  | CCGGGATTCC | GAGGAAGAAT  | GCACCGTAGG  | AGGGGCATGG | CCGGCCACGG |
| 17761 | CCTGACGGGC  | GGCATGCGTC | GTGCGCACCA  | CCGGCGGCGG  | CGCGCGTCGC | ACCGTCGCAT |
| 17821 | GCGCGGCGGT  | ATCCTGCCCC | TCCTTATTCC  | ACTGATCGCC  | GCGGCGATTG | GCGCCGTGCC |
| 17881 | CGGAATTGCA  | TCCGTGGCCT | TGCAGGCGCA  | GAGACACTGA  | TTAAAAACAA | GTTACATGTG |
| 17941 | GAAAAATCAA  | AATAAAAGTC | TGGACTCTCA  | CGCTCGCTTG  | GTCTGTAAAC | TATTTTGTAG |
| 18001 | AATGGAAGAC  | ATCAACTTTG | CGTCACTGGC  | CCCGCGACAC  | GGCTCGCGCC | CGTTCATGGG |
| 18061 | AAACTGGCAA  | GATATCGGCA | CCAGCAATAT  | GAGCGGTGGC  | GCCTTCAGCT | GGGGCTCGCT |
| 18121 | GTGGAGCGGC  | ATTAAAAATT | TCGGTTCCGC  | CGTTAAGAAC  | TATGGCAGCA | AAGCCTGGAA |
| 18181 | CAGCAGCACA  | GGCCAGATGC | TGAGGGACAA  | GTTGAAAGAG  | CAAAATTTCC | AACAAAAGGT |
| 18241 | GGTAGATGGC  | CTGGCCTCTG | GCATTAGCGG  | GGTGGTGGAC  | CTGGCCAACC | AGGCAGTGCA |
| 18301 | AAATAAGATT  | AACAGTAAGC | TTGATCCCCG  | CCCTCCCGTA  | GAGGAGCCTC | CACCGGCCGT |
| 18361 | GGAGACAGTG  | TCTCCAGAGG | GGCGTGGCGA  | AAAGCGTCCG  | CGACCCGACA | GGGAAGAAAC |
| 18421 | TCTGGTGACG  | CAAATAGACG | AGCCTCCCTC  | GTACGAGGAG  | GCACTAAAGC | AAGGCCTGCC |
| 18481 | CACCACCCGT  | CCCATCGCGC | CCATGGCTAC  | CGGAGTGCTG  | GGCCAGCACA | CACCCGTAAC |
| 18541 | GCTGGACCTG  | CCTCCCCCCG | CCGACACCCA  | GCAGAAACCT  | GTGCTGCCAG | GCCCGTCCGC |
| 18601 | CGTTGTTGTA  | ACCCGTCCTA | GCCGCGCGTC  | CCTGCGCCGC  | GCCGCCAGCG | GTCCGCGATC |
| 18661 | GTTGCGGCCC  | GTAGCCAGTG | GCAACTGGCA  | AAGCACACTG  | AACAGCATCG | TGGGTTTGGG |
| 18721 | GGTGCAATCC  | CTGAAGCGCC | GACGATGCTT  | CTGATAGCTA  | ACGTGTCGTA | TGTGTGTCAT |
| 18781 | GTATGCGTCC  | ATGTCGCCGC | CAGAGGAGCT  | GCTGAGCCGC  | CGCGCGCCCG | CTTTCCAAGA |
| 18841 | TGGCTACCCC  | TTCGATGATG | CCGCAGTGGT  | CTTACATGCA  | CATCTCGGGC | CAGGACGCCT |

FIG.11A-7

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|       |             |            |            |            |            |            |
|-------|-------------|------------|------------|------------|------------|------------|
| 18901 | CGGAGTACCT  | GAGCCCCGGG | CTGGTGCAGT | TCGCCCCGCG | CACCGAGACG | TACTTCAGCC |
| 18961 | TGAATAACAA  | GTTTAGAAAC | CCCACGGTGG | CGCCTACGCA | CGACGTGACC | ACAGACCGGT |
| 19021 | CTCAGCGTTT  | GACGCTGCGG | TTCATCCCCG | TGGACCGCGA | GGATACTGCG | TACTCGTACA |
| 19081 | AGGCGCGGTT  | CACCCTAGCT | GTGGGTGATA | ACCGTGTGCT | AGACATGGCT | TCCACGTACT |
| 19141 | TTGACATCCG  | CGGCGTGCTG | GACAGGGGCC | CTACTTTTAA | GCCCTACTCT | GGCACTGCCT |
| 19201 | ACAACGCACT  | GGCCCCCAAG | GGTGCCCCCA | ACTCGTGCGA | GTGGGAACAA | AATGAAACTG |
| 19261 | CACAAGTGGA  | TGCTCAAGAA | CTTGACGAAG | AGGAGAATGA | AGCCAATGAA | GCTCAGGCGC |
| 19321 | GAGAACAGGA  | ACAAGCTAAG | AAAACCCATG | TATATGCCCA | GGCTCCACTG | TCCGGAATAA |
| 19381 | AAATAACTAA  | AGAAGGTCTA | CAAATAGGAA | CTGCCGACGC | CACAGTAGCA | GGTGCCGGCA |
| 19441 | AAGAAATTTT  | CGCAGACAAA | ACTTTTCAAC | CTGAACCACA | AGTAGGAGAA | TCTCAATGGA |
| 19501 | ACGAAGCGGA  | TGCCACAGCA | GCTGGTGGA  | GGGTTCTTAA | AAAGACAAC  | CCCATGAAAC |
| 19561 | CCTGCTATGG  | CTCATACGCT | AGACCCACCA | ATTCCAACGG | CGGACAGGGC | GTTATGGTTG |
| 19621 | AACAAAATGG  | TAAATTGGAA | AGTCAAGTCG | AAATGCAATT | TTTTTCCACA | TCCACAAATG |
| 19681 | CCACAAATGA  | AGTTAACAAT | ATACAACCAA | CAGTTGTATT | GTACAGCGAA | GATGTAAACA |
| 19741 | TGGAAACTCC  | AGATACTCAT | CTTTCTTATA | AACCTAAAAT | GGGGGATAAA | AATGCCAAAG |
| 19801 | TCATGCTTGG  | ACAACAAGCA | ATGCCAAACA | GACCAAATTA | CATTGCTTTT | AGAGACAATT |
| 19861 | TTATTGGTCT  | CATGTATTAC | AACAGCACAG | GTAACATGGG | TGTCCTTGCT | GGTCAGGCAT |
| 19921 | CGCAGTTGAA  | CGCTGTTGTA | GATTTGCAAG | ACAGAAACAC | AGAGCTGTCC | TACCAGCTTT |
| 19981 | TGCTTGATTG  | AATTGGCGAC | AGAACAAGAT | ACTTTTCAAT | GTGGAATCAA | GCTGTTGACA |
| 20041 | GCTATGATCC  | AGATGTCAGA | ATTATTGAGA | ACCATGGAAC | TGAGGATGAG | TTGCCAAATT |
| 20101 | ATTGCTTTCC  | TCTTGGTGGA | ATTGGGATTA | CTGACACTTT | TCAAGCTGTT | AAAACAAC   |
| 20161 | CTGCTAACGG  | GGACCAAGGC | AATACTACCT | GGCAAAAAGA | TTCAACATTT | GCAGAACGCA |
| 20221 | ATGAAATAGG  | GGTGGGAAAT | AACTTTGCCA | TGGAAATTAA | CCTGAATGCC | AACCTATGGA |
| 20281 | GAAATTTTCT  | TTACTCCAAT | ATTGCGCTGT | ACCTGCCAGA | CAAGCTAAAA | TACAACCCCA |
| 20341 | CCAATGTGGA  | AATATCTGAC | AACCCCAACA | CCTACGACTA | CATGAACAAG | CGAGTGGTGG |
| 20401 | CTCCTGGGCT  | TGTAGACTGC | TACATTAACC | TTGGGGCGCG | CTGGTCTCTG | GACTACATGG |
| 20461 | ACAACGTAA   | TCCCTTTAAC | CACCACCGCA | ATGCGGGCCT | GCGTTACCGC | TCCATGTTGT |
| 20521 | TGGGAAACGG  | CCGCTACGTG | CCCTTTCACA | TTCAGGTGCC | CCAAAAGTTT | TTTGCCATTA |
| 20581 | AAAACCTCCT  | CCTCCTGCCA | GGCTCATACA | CATATGAATG | GAACCTCAGG | AAGGATGTTA |
| 20641 | ACATGGTTCT  | GCAGAGCTCT | CTGGGAAACG | ACCTTAGAGT | TGACGGGGCT | AGCATTAAGT |
| 20701 | TTGACAGCAT  | TTGTCTTTAC | GCCACCTTCT | TCCCCATGGC | CCACAACACG | GCCTCCACGC |
| 20761 | TGGAAGCCAT  | GCTCAGAAAT | GACACCAACG | ACCAGTCCTT | TAATGACTAC | CTTTCCGCCG |
| 20821 | CCAACATGCT  | ATATCCCAT  | CCCGCCAACG | CCACCAACGT | GCCCATCTCC | ATCCCATCGC |
| 20881 | GCAACTGGGC  | AGCATTTCGC | GGTTGGGCCT | TCACACGCTT | GAAGACAAAG | GAAACCCCTT |
| 20941 | CCCTGGGATC  | AGGCTACGAC | CCTTACTACA | CCTACTCTGG | CTCCATACCA | TACCTTGACG |
| 21001 | GAACCTTCTA  | TCTTAATCAC | ACCTTTAAGA | AGGTGGCCAT | TACTTTTGAC | TCTTCTGTTA |
| 21061 | GCTGGCCGGG  | CAACGACCGC | CTGCTTACTC | CCAATGAGTT | TGAGATTAAG | CGCTCAGTTG |
| 21121 | ACGGGGAGGG  | CTATAACGTA | GCTCAGTGCA | ACATGACAAA | GGACTGGTTC | CTAGTGCAGA |
| 21181 | TGTTGGCCAA  | CTACAATATT | GGCTACCAGG | GCTTCTACAT | TCCAGAAAGC | TACAAAGACC |
| 21241 | GCAATGTACTC | GTTCTTCAGA | AACTTCCAGC | CCATGAGCCG | GCAAGTGGTG | GACGATACTA |
| 21301 | AATACAAAGA  | TTATCAGCAG | GTTGGAATTA | TCCACCAGCA | TAACAAC    | GGCTTCGTAG |
| 21361 | GCTACCTCGC  | TCCCACCATG | CGCGAGGGAC | AAGCTTACCC | CGCTAATGTT | CCCTACCCAC |
| 21421 | TAATAGGCAA  | AACCGCGGTT | GATAGTATTA | CCCAGAAAAA | GTTTCTTTGC | GACCGACCC  |
| 21481 | TGTGGCGCAT  | CCCCTTCTCC | AGTAACTTTA | TGTCCATGGG | TGCGCTCACA | GACCTGGGCC |
| 21541 | AAAACCTTCT  | CTACGCAAA  | TCCGCCACG  | CGCTAGACAT | GACCTTTGAG | GTGGATCCCA |

FIG. 11A-8



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|       |            |             |            |             |            |            |
|-------|------------|-------------|------------|-------------|------------|------------|
| 21601 | TGGACGAGCC | CACCCCTTCTT | TATGTTTTGT | TTGAAGTCTT  | TGACGTGGTC | CGTGTGCACC |
| 21661 | AGCCGCACCG | CGGCGTCATC  | GAGACCGTGT | ACCTGCGCAC  | GCCCTTCTCG | GCCGGCAACG |
| 21721 | CCACAACATA | AAGAAGCAAG  | CAACATCAAC | AACAGCTGCC  | GCCATGGGCT | CCAGTGAGCA |
| 21781 | GGAAGTGAAG | GCCATTGTCA  | AAGATCTTGG | TTGTGGGCCA  | TATTTTTTGG | GCACCTATGA |
| 21841 | CAAGCGCTTC | CCAGGCTTTG  | TTTCCCCACA | CAAGCTCGCC  | TGCGCCATAG | TTAACACGGC |
| 21901 | CGGTGCGGAG | ACTGGGGGCG  | TACACTGGAT | GGCCTTTGCC  | TGGAACCCGC | GCTCAAAAAC |
| 21961 | ATGCTACCTC | TTTGAGCCCT  | TTGGCTTTTC | TGACCAACGT  | CTCAAGCAGG | TTTACCAGTT |
| 22021 | TGAGTACGAG | TCACTCCTGC  | GCCGTAGCGC | CATTGCCTCT  | TCCCCCGACC | GCTGTATAAC |
| 22081 | GCTGGAAAAG | TCCACCCAAA  | GCGTGCAGGG | GCCCAACTCG  | GCCGCCTGTG | GCCTATTCTG |
| 22141 | CTGCATGTTT | CTCCACGCCT  | TTGCCAACTG | GCCCCAAACT  | CCCATGGATC | ACAACCCAC  |
| 22201 | CATGAACCTT | ATTACCGGGG  | TACCCAACTC | CATGCTTAAC  | AGTCCCCAGG | TACAGCCAC  |
| 22261 | CCTGCGCCGC | AACCAGGAAC  | AGCTCTACAG | CTTCCTGGAG  | CGCCACTCGC | CCTACTTCCG |
| 22321 | CAGCCACAGT | GCGCAAATTA  | GGAGCGCCAC | TTCTTTTTGT  | CACTTGAAAA | ACATGTAAAA |
| 22381 | ATAATGTACT | AGGAGACACT  | TTCAATAAAG | GCAAATGTTT  | TTATTTGTAC | ACTCTCGGGT |
| 22441 | GATTATTTAC | CCCCACCCTT  | GCCGTCTGCG | CCGTTTAAAA  | ATCAAAGGGG | TTCTGCCGCG |
| 22501 | CATCGCTATG | CGCCACTGGC  | AGGGACACGT | TGCGATACTG  | GTGTTTAGTG | CTCCACTTAA |
| 22561 | ACTCAGGCAC | AACCATCCGC  | GGCAGCTCGG | TGAAGTTTTT  | ACTCCACAGG | CTGCGCACCA |
| 22621 | TCACCAACGC | GTTTAGCAGG  | TCGGGCGCCG | ATATCTTGAA  | GTCGCAGTTG | GGGCCTCCGC |
| 22681 | CCTGCGCGCG | CGAGTTGCGA  | TACACAGGGT | TACAGCACTG  | GAACACTATC | AGCGCCGGGT |
| 22741 | GGTGCACGCT | GGCCAGCACG  | CTCTTGTCGG | AGATCAGATC  | CGCGTCCAGG | TCCTCCGCGT |
| 22801 | TGCTCAGGGC | GAACGGAGTC  | AACCTTGGA  | GCTGCCTTCC  | CAAAAAGGGT | GCATGCCCAG |
| 22861 | GCTTTGAGTT | GCACTCGCAC  | CGTAGTGGCA | TCAGAAGGTG  | ACCGTGCCCA | GTCTGGGCGT |
| 22921 | TAGGATACAG | CGCCTGCATG  | AAAGCCTTGA | TCTGCTTAAA  | AGCCACCTGA | GCCTTTGCGC |
| 22981 | CTTCAGAGAA | GAACATGCCG  | CAAGACTTGC | CGGAAAACCTG | ATTGGCCGGA | CAGGCCGCGT |
| 23041 | CATGCACGCA | GCACCTTGCG  | TCGGTGTTGG | AGATCTGCAC  | CACATTTCCG | CCCCACCGGT |
| 23101 | TCTTCACGAT | CTTGGCCTTG  | CTAGACTGCT | CCTTCAGCGC  | GCGCTGCCCG | TTTTCGCTCG |
| 23161 | TCACATCCAT | TTCAATCACG  | TGCTCCTTAT | TTATCATAAT  | GCTCCCGTGT | AGACACTTAA |
| 23221 | GCTCGCCTTC | GATCTCAGCG  | CAGCGGTGCA | GCCACAACGC  | GCAGCCCGTG | GGCTCGTGGT |
| 23281 | GCTTGAGGT  | TACCTCTGCA  | AACGACTGCA | GGTACGCTG   | CAGGAATCGC | CCCATCATCG |
| 23341 | TCACAAAGGT | CTTGTTGCTG  | GTGAAGGTCA | GCTGCAACCC  | GCGGTGCTCC | TCGTTTAGCC |
| 23401 | AGGTCTTGCA | TACGGCCGCC  | AGAGCTTCCA | CTTGGTCAGG  | CAGTAGCTTG | AAGTTTGCCT |
| 23461 | TTAGATCGTT | ATCCACGTGG  | TACTTGTTCA | TCAACGCGCG  | CGCAGCCTCC | ATGCCCTTCT |
| 23521 | CCCACGCAGA | CACGATCGGC  | AGGCTCAGCG | GGTTTATCAC  | CGTGCTTTCA | CTTTCCGCTT |
| 23581 | CACTGGACTC | TTCTTTTCC   | TCTTGATCC  | GCATACCCCG  | CGCCACTGGG | TCGTCTTCAT |
| 23641 | TCAGCCGCCG | CACCGTGCGC  | TTACCTCCCT | TGCCGTGCTT  | GATTAGCACC | GGTGGGTTGC |
| 23701 | TGAAACCCAC | CATTTGTAGC  | GCCACATCTT | CTCTTTCTTC  | CTCGCTGTCC | ACGATCACCT |
| 23761 | CTGGGGATGG | CGGGCGCTCG  | GGCTTGGGAG | AGGGGCGCTT  | CTTTTTCTTT | TTGGACGCAA |
| 23821 | TGGCCAAATC | CGCCGTGCGAG | GTCGATGGCC | GCGGGCTGGG  | TGTGCGCGGC | ACCAGCGCAT |
| 23881 | CTTGTGACGA | GTCTTCTTCG  | TCCTCGGACT | CGAGACGCCG  | CCTCAGCCGC | TTTTTTGGGG |
| 23941 | GCGCGCGGGG | AGGCGGCGGC  | GACGGCGACG | GGGACGAGAC  | GTCCTCCATG | GTTGGTGGAC |
| 24001 | GTCGCGCCGC | ACCGCGTCCG  | CGCTCGGGGG | TGGTTTCGCG  | CTGCTCCTCT | TCCCGACTGG |
| 24061 | CCATTTCTTT | CTCCTATAGG  | CAGAAAAAGA | TCATGGAGTC  | AGTCGAGAAG | GAGGACAGCC |
| 24121 | TAACCGCCCC | CTTTGAGTTC  | GCCACCACCG | CCTCCACCGA  | TGCCGCCAAC | GCGCCTACCA |
| 24181 | CCTTCCCCGT | CGAGGCACCC  | CCGCTTGAGG | AGGAGGAAGT  | GATTATCGAG | CAGGACCCAG |
| 24241 | GTTTTGTAAG | CGAAGACGAC  | GAAGATCGCT | CAGTACCAAC  | AGAGGATAAA | AAGCAAGACC |

FIG. 11A-9

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|       |             |            |            |            |             |             |
|-------|-------------|------------|------------|------------|-------------|-------------|
| 24301 | AGGACGACGC  | AGAGGCAAAC | GAGGAACAAG | TCGGGCGGGG | GGACCAAAGG  | CATGGCGACT  |
| 24361 | ACCTAGATGT  | GGGAGACGAC | GTGCTGTTGA | AGCATCTGCA | GCGCCAGTGC  | GCCATTATCT  |
| 24421 | GCGACGCGTT  | GCAAGAGCGC | AGCGATGTGC | CCCTCGCCAT | AGCGGATGTC  | AGCCTTGCCCT |
| 24481 | ACGAACGCCA  | CCTGTTCTCA | CCGCGCGTAC | CCCCCAAACG | CCAAGAAAAC  | GGCACATGCG  |
| 24541 | AGCCCAACCC  | GCGCCTCAAC | TTCTACCCCG | TATTTGCCGT | GCCAGAGGTG  | CTTGCCACCT  |
| 24601 | ATCACATCTT  | TTTCCAAAAC | TGCAAGATAC | CCCTATCCTG | CCGTGCCAAC  | CGCAGCCGAG  |
| 24661 | CGGACAAGCA  | GCTGGCCTTG | CGGCAGGGCG | CTGTCATACC | TGATATCGCC  | TCGCTCGACG  |
| 24721 | AAGTGCCAAA  | AATCTTTGAG | GGTCTTGAGC | GCGACGAGAA | GCGCGCGGCA  | AACGCTCTGC  |
| 24781 | AACAAGAAAA  | CAGCGAAAAT | GAAAGTCACT | GTGGAGTGCT | GGTGGAACCT  | GAGGGTGACA  |
| 24841 | ACGCGCGCCT  | AGCCGTGCTG | AAACGCAGCA | TCGAGGTCAC | CCACTTTGCC  | TACCCGGCAC  |
| 24901 | TTAACCTACC  | CCCCAAGGTT | ATGAGCACAG | TCATGAGCGA | GCTGATCGTG  | CGCCGTGCAC  |
| 24961 | GACCCCTGGA  | GAGGGATGCA | AACTTGCAAG | AACAAACCGA | GGAGGGCCTA  | CCCGCAGTTG  |
| 25021 | GCGATGAGCA  | GCTGGCGCGC | TGGCTTGAGA | CGCGCGAGCC | TGCCGACTTG  | GAGGAGCGAC  |
| 25081 | GCAAGCTAAT  | GATGGCCGCA | GTGCTTGTTA | CCGTGGAGCT | TGAGTGCATG  | CAGCGGTTCT  |
| 25141 | TTGCTGACCC  | GGAGATGCAG | CGCAAGCTAG | AGGAAACGTT | GCACTACACC  | TTTCGCCAGG  |
| 25201 | GCTACGTGCG  | CCAGGCCTGC | AAAATTTCCA | ACGTGGAGCT | CTGCAACCTG  | GTCTCCTACC  |
| 25261 | TTGGAATTTT  | GCACGAAAAC | CGCCTTGGGC | AAAACGTGCT | TCATTCCACG  | CTCAAGGGCG  |
| 25321 | AGGCGCGCCG  | CGACTACGTC | CGCGACTGCG | TTTACTTATT | TCTGTGCTAC  | ACCTGGCAAA  |
| 25381 | CGGCCATGGG  | CGTGTGGCAG | CAGTGCCTGG | AGGAGCGCAA | CCTGAAGGAG  | CTGCAGAAGC  |
| 25441 | TGCTAAAGCA  | AAACTTGAAG | GACCTATGGA | CGGCCTTCAA | CGAGCGCTCC  | GTGGCCGCGC  |
| 25501 | ACCTGGCGGA  | CATTATCTTC | CCCGAACGCC | TGCTTAAAAC | CCTGCAACAG  | GGTCTGCCAG  |
| 25561 | ACTTCACCAG  | TCAAAGCATG | TTGCAAACT  | TTAGGAACTT | TATCCTAGAG  | CGTTCAGGAA  |
| 25621 | TTCTGCCCGC  | CACCTGCTGT | GCGCTTCCTA | GCGACTTTGT | GCCCATTAA   | TACCGTGAAT  |
| 25681 | GCCCTCCGCC  | GCTTTGGGGT | CACTGCTACC | TTCTGCAGCT | AGCCAACTAC  | CTTGCCCTACC |
| 25741 | ACTCCGACAT  | CATGGAAGAC | GTGAGCGGTG | ACGGCCTACT | GGAGTGTAC   | TGTCGCTGCA  |
| 25801 | ACCTATGCAC  | CCCGCACCGC | TCCCTGGTCT | GCAATTCACA | ACTGCTTAGC  | GAAAGTCAAA  |
| 25861 | TTATCGGTAC  | CTTTGAGCTG | CAGGGTCCCT | CGCCTGACGA | AAAGTCCGCG  | GCTCCGGGGT  |
| 25921 | TGAAACTCAC  | TCCGGGGCTG | TGGACGTCGG | CTTACCTTCG | CAAAATTTGTA | CCTGAGGACT  |
| 25981 | ACCACGCCCA  | CGAGATTAGG | TTCTACGAAG | ACCAATCCCG | CCCGCCAAAT  | GCGGAGCTTA  |
| 26041 | CCGCCTGCGT  | CATTACCCAG | GGCCACATCC | TTGGCCAATT | GCAAGCCATT  | AACAAAGCCC  |
| 26101 | GCCAAGAGTT  | TCTGCTACGA | AAGGGACGGG | GGGTTTACTT | GGACCCCCAG  | TCCGGCGAGG  |
| 26161 | AGCTCAACCC  | AATCCCCCGG | CCGCCGCAGC | CCTATCAGCA | GCCGCGGGCC  | CTTGCTTCCC  |
| 26221 | AGGATGGCAC  | CCAAAAAGAA | GCTGCAGCTG | CCGCCGCCGC | CACCCACGGA  | CGAGGAGGAA  |
| 26281 | TACTGGGACA  | GTCAGGCAGA | GGAGGTTTTG | GACGAGGAGG | AGGAGATGAT  | GGAAGACTGG  |
| 26341 | GACAGCCTAG  | ACGAGGAAGC | TTCCGAGGCC | GAAGAGGTGT | CAGACGAAAC  | ACCGTCACCC  |
| 26401 | TCGGTGCAT   | TCCCCTCGCC | GGCGCCCCAG | AAATCGGCAA | CCGTTCCCAG  | CATTGCTACA  |
| 26461 | ACCTCCGCTC  | CTCAGGCGCC | GCCGGCACTG | CCCGTTCCGC | GACCCAACCG  | TAGATGGGAC  |
| 26521 | ACCACTGGAA  | CCAGGGCCGG | TAAGTCTAAG | CAGCCGCCGC | CGTTAGCCCA  | AGAGCAACAA  |
| 26581 | CAGCGCCAAG  | GCTACCGCTC | GTGGCGCGTG | CACAAGAACG | CCATAGTTGC  | TTGCTTGCAA  |
| 26641 | GA CTGTGGGG | GCAACATCTC | CTTCGCCCGC | CGTTTTCTTC | TCTACCATCA  | CGGCGTGGCC  |
| 26701 | TTCCCCCGTA  | ACATCCTGCA | TTACTACCGT | CATCTCTACA | GCCCCTACTG  | CACCGGCGGC  |
| 26761 | AGCGGCAGCA  | ACAGCAGCGG | CCACGCAGAA | GCAAAGGCGA | CCGGATAGCA  | AGACTCTGAC  |
| 26821 | AAAGCCCAAG  | AAATCCACAG | CGGCGGCAGC | AGCAGGAGGA | GGAGCACTGC  | GTCTGGCGCC  |
| 26881 | CAACGAACCC  | GTATCGACCC | GCGAGCTTAG | AAACAGGATT | TTTCCCACTC  | TGTATGCTAT  |
| 26941 | ATTTCAACAG  | AGCAGGGGCC | AAGAACAAGA | GCTGAAAATA | AAAAACAGGT  | CTCTGCGCTC  |

FIG.11A-10



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|       |            |             |            |            |            |             |
|-------|------------|-------------|------------|------------|------------|-------------|
| 27001 | CCTCACCCGC | AGCTGCCTGT  | ATCACAAAAG | CGAAGATCAG | CTTCGGCGCA | CGCTGGAAGA  |
| 27061 | CGCGGAGGCT | CTCTTCAGCA  | AATACTGCGC | GCTGACTCTT | AAGGACTAGT | TTCGCGCCCT  |
| 27121 | TTCTCAAATT | TAAGCGCGAA  | AACTACGTCA | TCTCCAGCGG | CCACACCCGG | CGCCAGCACC  |
| 27181 | TGTCGTCAGC | GCCATTATGA  | GCAAGGAAAT | TCCCACGCCC | TACATGTGGA | GTTACCAGCC  |
| 27241 | ACAAATGGGA | CTTGCGGCTG  | GAGCTGCCCA | AGACTACTCA | ACCCGAATAA | ACTACATGAG  |
| 27301 | CGCGGGACCC | CACATGATAT  | CCCGGGTCAA | CGGAATCCGC | GCCCACCGAA | ACCGAATTCT  |
| 27361 | CCTCGAACAG | GCGGCTATTA  | CCACCACACC | TCGTAATAAC | CTTAATCCCC | GTAGTTGGCC  |
| 27421 | CGCTGCCCTG | GTGTACCAGG  | AAAGTCCCGC | TCCCACCACT | GTGGTACTTC | CCAGAGACGC  |
| 27481 | CCAGGCCGAA | GTTCAGATGA  | CTAACTCAGG | GGCGCAGCTT | GCGGGCGGCT | TTCGTACACAG |
| 27541 | GGTGCGGTCG | CCCGGGCAGG  | GTATAACTCA | CCTGAAAATC | AGAGGGCGAG | GTATTACAGCT |
| 27601 | CAACGACGAG | TCGGTGAGCT  | CCTCTCTTGG | TCTCCGTCCG | GACGGGACAT | TTCAGATCGG  |
| 27661 | CGGCGCTGGC | CGCTCTTCAT  | TTACGCCCCG | TCAGGCGATC | CTAACTCTGC | AGACCTCGTC  |
| 27721 | CTCGGAGCCG | CGCTCCGGAG  | GCATTGGAAC | TCTACAATTT | ATTGAGGAGT | TCGTGCCTTC  |
| 27781 | GGTTTACTTC | AACCCCTTTT  | CTGGACCTCC | CGGCCACTAC | CCGGACCACT | TTATTCCCAA  |
| 27841 | CTTTGACGCG | GTAAAAGACT  | CGGCGGACGG | CTACGACTGA | ATGACCACTG | GAGAGGCAGA  |
| 27901 | GCAACTGCGC | CTGACACACC  | TCGACCACTG | CCGCCGCCAC | AAGTGCTTTG | CCGCGGGCTC  |
| 27961 | CGGTGAGTTT | TGTTACTTTG  | AATTGCCCGA | AGAGCATATC | GAGGGCCCGG | CGCACGGCGT  |
| 28021 | CCGGCTCACC | ACCCAGGTAG  | AGCTTACACG | TAGCCTGATT | CGGGAGTTTA | CCAAGCGCCC  |
| 28081 | CCTGCTAGTG | GAGCGGGAGC  | GGGGTCCCTG | TGTTCTGACC | GTGGTTTGCA | ACTGTCCTAA  |
| 28141 | CCCTGGATTA | CATCAAGATC  | TTTGTTGTCA | TCTCTGTGCT | GAGTATAATA | AATACAGAAA  |
| 28201 | TTAGAATCTA | CTGGGGCTCC  | TGTCGCCATC | CTGTGAACGC | CACCGTTTTT | ACCCACCCAA  |
| 28261 | AGCAGACCAA | AGCAAACCTC  | ACCTCCGGTT | TGCACAAGCG | GGCCAATAAG | TACCTTACCT  |
| 28321 | GGTACTTTAA | CGGCTCTTCA  | TTTGTAATTT | ACAACAGTTT | CCAGCGAGAC | GAAGTAAGTT  |
| 28381 | TGCCACACAA | CCTTCTCGGC  | TTCAACTACA | CCGTCAAGAA | AAACACCACC | ACCACCCTCC  |
| 28441 | TCACCTGCCG | GGAACGTACG  | AGTGCCTCAC | CGGTTGCTGC | GCCCACACCT | ACAGCCTGAG  |
| 28501 | CGTAACCAGA | CATTACTCCC  | ATTTTCCCAA | AACAGGAGGT | GAGCTCAACT | CCCGGAACTC  |
| 28561 | AGGTCAAAAA | AGCATTTTGC  | GGGGTGCTGG | GATTTTTTAA | TTAAGTATAT | GAGCAATTCA  |
| 28621 | AGTAACTCTA | CAAGCTTGTC  | TAATTTTTCT | GGAATTGGGG | TCGGGGTTAT | CCTTACTCTT  |
| 28681 | GTAATTCTGT | TTATTCTTAT  | ACTAGCACTT | CTGTGCCTTA | GGGTTGCCGC | CTGCTGCACG  |
| 28741 | CACGTTTGTA | CCTATTGTCA  | GCTTTTTAAA | CGCTGGGGGC | GACATCCAAG | ATGAGGTACA  |
| 28801 | TGATTTTAGG | CTTGCTCGCC  | CTTGCGGCAG | TCTGCAGCGC | TGCCAAAAAG | GTTGAGTTTA  |
| 28861 | AGGAACCAGC | TTGCAATGTT  | ACATTTAAAT | CAGAAGCTAA | TGAATGCACT | ACTCTTATAA  |
| 28921 | AATGCACCAC | AGAACATGAA  | AAGCTTATTA | TTCGCCACAA | AGACAAAATT | GGCAAGTATG  |
| 28981 | CTGTATATGC | TATTTGGCAG  | CCAGGTGACA | CTAACGACTA | TAATGTCACA | GTCTTCCAAG  |
| 29041 | GTGAAAATCG | TAAAACTTTT  | ATGTATAAAT | TTCCATTTTA | TGAAATGTGC | GATATTACCA  |
| 29101 | TGTACATGAG | CAAACAGTAC  | AAGTTGTGGC | CCCCACAAAA | GTGTTTAGAG | AACACTGGCA  |
| 29161 | CCTTTTGTTT | CACCGCTCTG  | CTTATTACAG | CGCTTGCTTT | GGTATGTACC | TTACTTTATC  |
| 29221 | TCAAATACAA | AAGCAGACGC  | AGTTTTATTG | ATGAAAAGAA | AATGCCTTGA | TTTTCCGCTT  |
| 29281 | GCTTGTATTC | CCCTGGACAA  | TTTACTCTAT | GTGGGATATG | CGCCAGGCGG | GAAAGATTAT  |
| 29341 | ACCCACAACC | TTCAAATCAA  | ACTTTCTCTG | ACGTTAGCGC | CTGACTTCTG | CCAGCGCCTG  |
| 29401 | CACTGCAAAT | TTGATCAAAC  | CCAGCTTCAG | CTTGCCTGCT | CCAGAGATGA | CCGGCTCAAC  |
| 29461 | CATCGCGCCC | ACAACGGACT  | ATCGCAACAC | CACTGCTACC | GGACTAAAAT | CTGCCCTAAA  |
| 29521 | TTTACCCCAA | GTTTCATGCCT | TTGTCAATGA | CTGGGCGAGC | TTGGGCATGT | GGTGGTTTTT  |
| 29581 | CATAGCGCTT | ATGTTTGTTT  | GCCTTATTAT | TATGTGGCTT | ATTTGTTGCC | TAAAGCGCAG  |
| 29641 | ACGCGCCAGA | CCCCCATCT   | ATAGGCCTAT | CATTGTGCTC | AACCCACACA | ATGAAAAAAT  |

FIG.11A-11

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29701 TCATAGATTG GACGGTCTCA AACCATGTTC TCTTCTTTTA CAGTATGATT AAATGAGACA  
29761 TGATTCCTCG AGTCCTTATA TTATTGACCC TTGTTGCGCT TTTCTGTGCG TGCTCTACAT  
29821 TGGCTGCGGT CGCTCACATC GAAGTAGATT GCATCCCACC TTTCACAGTT TACCTGCTTT  
29881 ACGGATTTGT CACCCTTATC CTCATCTGCA GCCTCGTCAC TGTAGTCATC GCCTTCATTC  
29941 AGTTCATTGA CTGGATTTGT GTGCGCATTG CGTACCTTAG GCACCATCCG CAATACAGAG  
30001 ACAGGACTAT AGCTGATCTT CTCAGAATTC TTTAATTATG AAACGGATTG TCACTTTTGT  
30061 TTTGCTGATT TTCTGCGCCC TACCTGTGCT TTGCTCCCAA ACCTCAGCGC CTCCCAAAAG  
30121 ACATATTTCC TGCAGATTCA CTCAAATATG GAACATTCCC AGCTGCTACA ACAAACAGAG  
30181 CGATTTGTCA GAAGCCTGGT TATACGCCAT CATCTCTGTC ATGGTTTTTT GCAGTACCAT  
30241 TTTTGCCCTA GCCATATAACC CATACCTTGA CATTGGTTGG AATGCCATAG ATGCCATGAA  
30301 CCACCCTACT TTCCCAGCGC CCAATGTCAT ACCACTGCAA CAGGTTATTG CCCCAATCAA  
30361 TCAGCCTCGC CCCCCTTCTC CCACCCCCAC TGAGATTAGC TACTTTAATT TGACAGGTGG  
30421 AGATGACTGA ATCTCTAGAT CTAGAATTGG ATGGAATTAA CACCGAACAG CGCCTACTAG  
30481 AAAGGCGCAA GGCGGCGTCC GAGCGAGAAC GCCTAAACA AGAAGTTGAA GACATGGTTA  
30541 ACCTGCACCA GTGTAAAAGA GGTATCTTTT GTGTGGTCAA GCAGGCCAAA CTTACCTACG  
30601 AAAAAACCAC TACCGGCAAC CGCCTTAGCT ACAAGCTACC CACCCAGCGC CAAAACTGG  
30661 TGCTTATGGT GGGAGAAAAA CCTATCACCG TCACCCAGCA CTCGGCAGAA ACAGAAGGCT  
30721 GCCTGCACTT CCCCTATCAG GGTCCAGAGG ACCTCTGCAC TCTTATTAAC ACCATGTGTG  
30781 GCATTAGAGA TCTTATTCCA TTCAACTAAC AATAAACACA CAATAAATTA CTTACTTAAA  
30841 ATCAGTCAGC AAATCTTTGT CCAGCTTATT CAGCATCACC TCCTTTCCCT CCTCCCAACT  
30901 CTGGTATTTT AGCAGCCTTT TAGCTGCGAA CTTTCTCCAA AGTCTAAATG GGATGTCAAA  
30961 TTCCTCATGT TCTTGCCCT CCGCACCCAC TATCTTCATA TTGTTGCAGA TGAAACGCGC  
31021 CAGACCGTCT GAAGACACCT TCAACCCTGT GTACCCATAT GACACGGAAC CCGGCCCTCC  
31081 AACTGTGCCT TTCCTTACCC CTCCCTTTGT GTCGCCAAAT GGGTTCCAAG AAAGTCCCCC  
31141 CGGAGTGCTT TCTTTGCGTC TTTCAGAACC TTTGGTTACC TCACACGGCA TGCTTGCGCT  
31201 AAAAATGGGC AGCGGCCTGT CCCTGGATCA GGCAGGCAAC CTTACATCAA ATACAATCAC  
31261 TGTTTCTCAA CCGCTAAAAA AAACAAAGTC CAATATAACT TTGGAAACAT CCGCGCCCT  
31321 TACAGTCAGC TCAGGCGCCC TAACCATGGC CACAACCTCG CCTTTGGTGG TCTCTGACAA  
31381 CACTCTTACC ATGCAATCAC AAGCACCGCT AACCCTGCAA GACTCAAAAC TTAGCATTGC  
31441 TACCAAAGAG CCACTTACAG TGTAGATGG AAAACTGGCC CTGCAGACAT CAGCCCCCT  
31501 CTCTGCCACT GATAACAACG CCCTCACTAT CACTGCCTCA CCTCCTCTTA CTACTGCAAA  
31561 TGGTAGTCTG GCTGTTACCA TGGAAAACCC ACTTTACAAC AACATGGAA AACTTGGGCT  
31621 CAAAATTGGC GGTCTTTTGC AAGTGGCCAC CGACTCACAT GCACTAACAC TAGGTACTGG  
31681 TCAGGGGGTT GCAGTTCATA ACAATTTGCT ACATACAAAA GTTACAGGCG CAATAGGGTT  
31741 TGATACATCT GGCAACATGG AACTTAAAAC TGGAGATGGC CTCTATGTGG ATAGCGCCGG  
31801 TCCTAACCAA AAATAACATA TTAATCTAAA TACCACAAAA GGCCTTGCTT TTGACAACAC  
31861 CGCAATAACA ATTAACGCTG GAAAAGGGTT GGAATTTGAA ACAGACTCCT CAAACGGAAC  
31921 TCCCATAAAA ACAAAAATTG GATCAGGCAT ACAATATAAT ACCAATGGAG CTATGGTTGC  
31981 AAAACTTGGA ACAGGCCTCA GTTTTGACAG CTCCGGAGCC ATAACAATGG GCAGCATAAA  
32041 CAATGACAGA CTTACTCTTT GGACAACACC AGACCCATCC CCAAATTGCA GAATTGCTTC  
32101 AGATAAAGAC TGCAAGCTAA CTCTGGCGCT AACAAAATGT GGCAGTCAAA TTTTGGGCAC  
32161 TGTTTCAGCT TTGGCAGTAT CAGGTAATAT GGCCTCCATC AATGGAATC TAAGCAGTGT  
32221 AAACCTGGTT CTTAGATTTG ATGACAACGG AGTGCTTATG TCAAATTCAT CACTGGACAA  
32281 ACAGTATTGG AACTTTAGAA ACGGGGACTC CACTAACGGT CAACCATACA CTTATGCTGT  
32341 TGGGTTTATG CCAAACCTAA AAGCTTACCC AAAAATCAA AGTAAAATG CAAAAAGTAA

FIG. 11A-12

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32401 TATTGTTAGC CAGGTGTATC TTAATGGTGA CAAGTCTAAA CCATTGCATT TTAATTATTAC  
32461 GCTAAATGGA ACAGATGAAA CCAACCAAGT AAGCAAATAC TCAATATCAT TCAGTTGGTC  
32521 CTGGAACAGT GGACAATACA CTAATGACAA ATTTGCCACC AATTCCTATA CCTTCTCCTA  
32581 CATTGCCCAG GAATAAAGAA TCGTGAACCT GTTGCATGTT ATGTTTCAAC GTGTTTATTT  
32641 TTCAATTGCA GAAAATTTCA AGTCATTTTT CATTCAGTAG TATAGCCCCA CCACCACATA  
32701 GCTTATACTA ATCACC GTAC CTTAATCAAA CTCACAGAAC CCTAGTATTC AACCTGCCAC  
32761 CTCCCTCCCA ACACACAGAG TACACAGTCC TTTCTCCCCG GCTGGCCTTA AACAGCATCA  
32821 TATCATGGGT AACAGACATA TTCTTAGGTG TTATATTCCA CACGGTCTCC TGTCGAGCCA  
32881 AACGCTCATC AGTGATGTTA ATAACTCCC CGGGCAGCTC GCTTAAGTTC ATGTCGCTGT  
32941 CCAGCTGCTG AGCCACAGGC TGCTGTCCAA CTTGCGGTTG CTCAACGGGC GGCGAAGGAG  
33001 AAGTCCACGC CTACATGGGG GTAGAGTCAT AATCGTGCAT CAGGATAGGG CGGTGGTGCT  
33061 GCAGCAGCGC GCGAATAAAC TGCTGCCGCC GCCGCTCCGT CCTGCAGGAA TACAACATGG  
33121 CAGTGGTCTC CTCAGCGATG ATTCGCACCG CCCGCAGCAT AAGGCGCCTT GTCCTCCGGG  
33181 CACAGCAGCG CACCCTGATC TCACTTAAGT CAGCACAGTA ACTGCAGCAC AGTACCACAA  
33241 TATTGTTTAA AATCCACAG TGCAAGGCGC TGTATCCAAA GCTCATGGCG GGGACCACAG  
33301 AACCCACGTG GCCATCATAC CACAAGCGCA GGTAGATTAA GTGGCGACCC CTCATAAACA  
33361 CGCTGGACAT AAACATTACC TCTTTTGGCA TGTTGTAATT CACCACCTCC CGGTACCATA  
33421 TAAACCTCTG ATTAACATG GCGCCATCCA CCACCATCCT AAACCAGCTG GCCAAAACCT  
33481 GCCCGCCGGC TATGCACTGC AGGGAACCGG GACTGGAACA ATGACAGTGG AGAGCCCAGG  
33541 ACTCGTAACC ATGGATCATC ATGCTCGTCA TGATATCAAT GTTGGCACAA CACAGGCACA  
33601 CGTGCATACA CTTCTCAGG ATTACAAGCT CCTCCCGCT CAGAACCATA TCCCAGGGAA  
33661 CAACCCATTC CTGAATCAGC GTAAATCCCA CACTGCAGGG AAGACCTCGC ACGTAACTCA  
33721 CGTTGTGCAT TGTCAAAGTG TTACATTCGG GCAGCAGCGG ATGATCCTCC AGTATGGTAG  
33781 CGCGTGTCTC TGTCTCAAAA GGAGGTAGGC GATCCCTACT GTACGGAGTG CGCCGAGACA  
33841 ACCGAGATCG TGTTGGTCGT AGTGTCTATGC CAAATGGAAC GCCGGACGTA GTCATATTTC  
33901 CTGAAGCAAA ACCAGGTGCG GCGGTGACAA ACAGATCTGC GTCTCCGGTC TCGTCGCTTA  
33961 GCTCGCTCTG TGTAAGTAGT GTAGTATATC CACTCTCTCA AAGCATCCAG GCGCCCCCTG  
34021 GCTTCGGGTT CTATGTAAAC TCCTTCATGC GCCGCTGCC TGATAACATC CACCACCGCA  
34081 GAATAAGCCA CACCCAGCCA ACCTACACAT TCGTTCTGCG AGTCACACAC GGGAGGAGCG  
34141 GGAAGAGCTG GAAGAACCAT GTTTTTTTTT TTTATTCCAA AAGATTATCC AAAACCTCAA  
34201 AATGAAGATC TATTAAGTGA ACGCGCTCCC CTCCGGTGGC GTGGTCAAAC TCTACAGCCA  
34261 AAGAACAGAT AATGGCATT GTAGATGTT GCACAATGGC TTCCAAAAGG CAAACTGCCC  
34321 TCACGTCCAA GTGGACGTAA AGGCTAAACC CTTCAGGGTG AATCTCCTCT ATAAACATTC  
34381 CAGCACCTTC AACCATGCCC AAATAATTTT CATCTCGCCA CTTATCAAT ATGTCTCTAA  
34441 GCAAATCCCG AATATTAAGT CCGGCCATTG TAAAAATCTG CTCCAGAGCG CCCTCCACCT  
34501 TCAGCCTCAA GCAGCGAATC ATGATTGCAA AAATTCAGGT TCCTCACAGA CCTGTATAAG  
34561 ATTCAAAAGC GGAACATTAA CAAAAATACC GCGATCCCGT AGGTCCCTTC GCAGGGCCAG  
34621 CTGAACATAA TCGTGCAGGT CTGCACGGAC CAGCGCGGCC ACTTCCCCGC CAGGAACCAT  
34681 GACAAAAGAA CCCCACTGA TTATGACACG CATACTCGGA GCTATGCTAA CCAGCGTAGC  
34741 CCCGATGTAA GCTTGTTGCA TGGGCGGCGA TATAAAATGC AAGGTACTGC TCAAAAAATC  
34801 AGGCAAAGCC TCGCGCAAAA AAGCAAGCAC ATCGTAGTCA TGCTCATGCA GATAAAGGCA  
34861 GGTAAGTTCC GGAACCACCA CAGAAAAAGA CACCATTTTT CTCTCAAACA TGTCTGCGGG  
34921 TTCCTGCATA AACACAAAAT AAAATAACAA AAAAAAAAAA ACATTTAAAC ATTAGAAGCC  
34981 TGTNTTACAA CAGGAAAAAC AACCCTTATA AGCATAAGAC GGACTACGGC CATGCCGGCG  
35041 TGACCGTAAA AAAACTGGTC ACCGTGATTA AAAAGCACCA CCGACAGTTC CTCGGTCTAG

FIG. 11A-13

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35101 TCCGGAGTCA TAATGTAAGA CTCGGTAAAC ACATCAGGTT GGTTAACATC GGTCAGTGCT
35161 AAAAAGCGAC CGAAATAGCC CGGGGGAATA CATACCCGCA GCGGTAGAGA CAACATTACA
35221 GCCCCCATAG GAGGTATAAC AAAATTAATA GGAGAGAAAA ACACATAAAC ACCTGAAAAA
35281 CCCTCCTGCC TAGGCAAAAT AGCACCTCC CGCTCCAGAA CAACATACAG CGCTTCCACA
35341 GCGGCAGCCA TAACAGTCAG CTTACCAGT AAAAAACCT ATTAAAAAAC ACCACTCGAC
35401 ACGGCACCAG CTCAATCAGT CACAGTGTA AAAGGGCCAA GTACAGAGCG AGTATATATA
35461 GGAATAAAAA ATGACGTAAC GGTAAAGTC CAAAAAAC ACCCAGAAAA CCGCACGCGA
35521 ACCTACGCCC AGAAACGAAA GCCAAAAAAC CCACAACTTC CTCAAATCTT CACTTCCGTT
35581 TTCCCACGAT ACGTCACTTC CCATTTTAAA AAAAACTAC AATTCCCAAT ACATGCAAGT
35641 TACTCCGCCC TAAACCTAC GTCACCCGCC CCGTTCCAC GCCCCGCGCC ACGTCACAAA
35701 CTCCACCCC TCATTATCAT ATGGCTTCA ATCCAAAATA AGGTATATTA TTGATGATG
```

FIG.11A-14

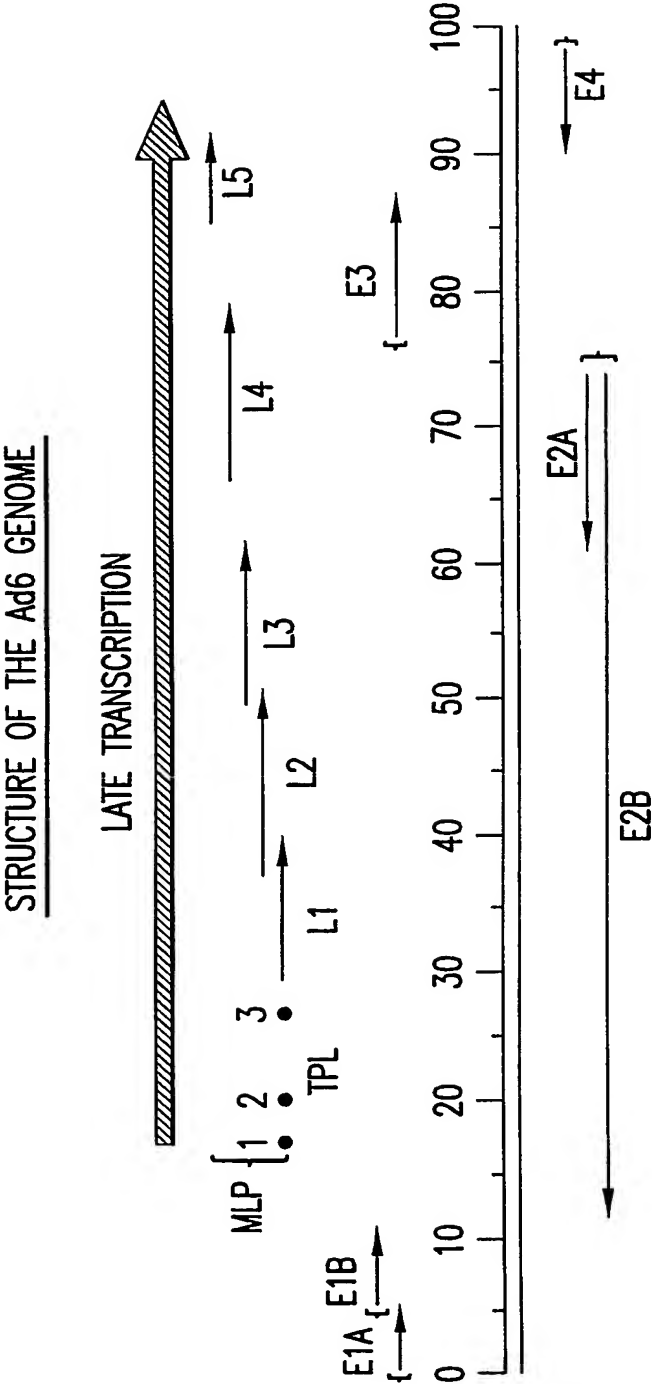


FIG.12

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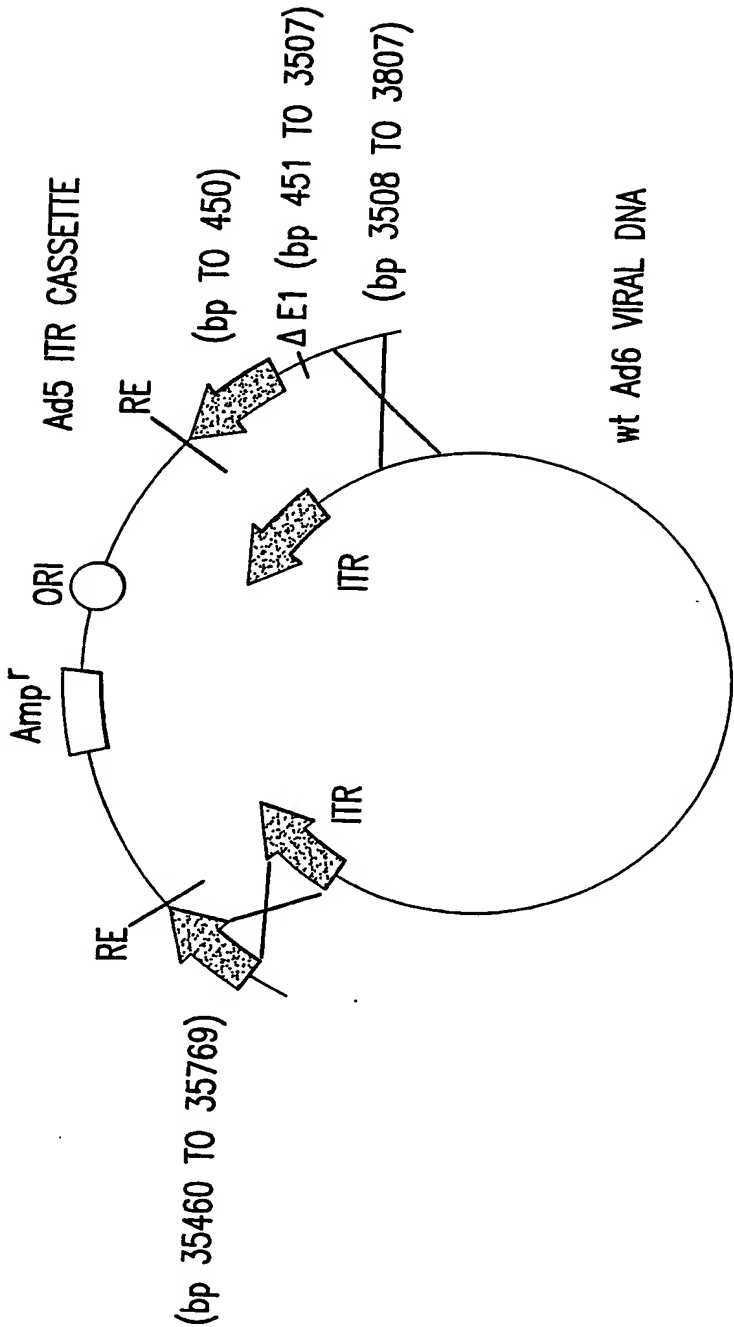


FIG.13